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ALIGNMENTS

RESULT 1
ADM16603
ID ADM1 WPI; 2004-270022/25. N-PSDB; ADM16606. Cytostatic; nootropic; neuroprotective; antiparkinsonian; inhibitor; cell migration; cell invasion; cell proliferation; angiogenesis; apoptosis; tumour growth rate; human kininogen domain 3v; HK-D3v; HK neurodegenerative disease; Alzheimer's disease; Parkinson's disease; Donate 13-SEP-2002; 2002US-0410279P 15-SEP-2003; 2003WO-US028881. Synthetic. 25-MAR-2004. WO2004024877-A2 Homo sapiens. amyotrophic lateral sclerosis; ALS; mutein. HK-D3v(GS) 17-JUN-2004 ADM16603; ADM16603 standard; protein; 125 (ATTE-) ATTENUON LLC 'n Mazar amino acid sequence, (first entry) A, веq ₽ id Ŋ..

HK-D3;

Novel isolated anti-angiogenic polypeptide comprising sequence of human kiningen domain 3v (HK-D3v) or variant of native HK-D3, useful for inhibiting angiogenesis, endothelial cell proliferation or endothelial tube formation.

Disclosure; SEQ ID NO 2; 79pp; English.

The invention relates to an isolated anti-angiogenic polypeptide (I) comprising the sequence of human kininogen domain 3v (HK-D3v) (fully defined sequence (S1) of 127 amino acids as given in specification) or variant or derivative of native HK-D3 (fully defined sequence (S2) of 123 amino acids as given in specification), and has at least 20% of activity of native HK-D3 in inhibiting angiogenesis, endothelial cell

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CC diagnostically or therapeutically labelled anti-angiogenic polypeptide (II) comprising (I) labelled with a diagnostic or therapeutic label, and CC (II) comprising (I) labelled with a diagnostic or therapeutic label, and CC with a detectable label and a carrier. Further disclosed is an affinity CC ligand (IX) useful for binding to or isolating an HK-D3-binding molecule CC or cells expressing the binding molecule, comprising (II) immobilized to a correction, cell invasion, cell proliferation or angiogenesis, or for cells expressing which involves contacting cells associated with CC inducing apoptosis which involves contacting cells associated with CC undesired cell migration, invasion, proliferation or angiogenesis with CC (I) is useful for treating neurodegenerative diseases e.g., CC (IX). The current sequence represents an N-terminal addition variant of HK-D3 designated HD-D3v(GS), that includes an additional GS at the N-terminal columns of the N-terminal additional CS at the N-terminal addition variant of terminal columns of the N-terminal additional CS at the N-terminal cS at the N-terminal additional CS at the N-terminal cS at the N-terminal additional CS at the N-terminal cS at the N-terminal additional CS at the N-terminal cS
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cancer; cytostatic.
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                   Novel human kininogen domain 3 variant inhibiting angiogenesis, endothelial cell proliferation or endothelial tube formation, utreating disease associated with undesired cell migration, inveproliferation or angiogenesis.
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Pred. No. 3.3e-67;
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                                                                                                                                                                                                                                                                                                                                                                                                                 Anti-angiogenic; angiogenesis; inhibitor; kininogen; homologue; endothelial cell proliferation; apoptosis; cancer; ocular disorrheumatoid arthritis; cytostatic; antiarthritic; antirheumatic; human; D3 peptide.
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Matches 123
Novel isolated anti-angiogenic polypeptide comprising sequence of human kiningen domain 3v (HK-D3v) or variant of native HK-D3, useful for
                                                                                                 N-PSDB; ADM16605.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure;
                                                                                                                         2004-270022/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present sequence is that of domain 3 of human high mol.wt.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    amino acid sequence,
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HK-D3v; HK-D3;
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cc variant or derivative of native HK-D3 (fully defined sequence (S2) of 123 cc amino acids as given in specification), and has at least 20% of activity of native HK-D3 in inhibiting angiogenesis, endothelial cell conformation. Also disclosed is a comprisit or therapeutically labelled anti-angiogenic polypeptide comprisiting (I) labelled with a diagnostic or therapeutic label, and composition (II) comprising (I) labelled with a diagnostic or therapeutic label, and composition (II) comprising (II) labelled with a diagnostic or therapeutic label, and composition (II) comprising (II) labelled composition (II) comprising (II) abelled with a detectable label and a carrier. Further disclosed is an affinity ligand (IX) useful for binding to or isolating an HK-D3-binding molecule concells expressing the binding molecule, comprising (I) immobilized to a solid support or carrier. (I) or (II) are useful for inhibiting cell migration, cell invasion, cell proliferation or anglogenesis, or for undesired cell migration, invasion, proliferation or anglogenesis with
Sequence 123 AA;

    (1) is useful for treating neurodegenerative diseases e.g.,
Alzheimer's disease, Parkinson's disease or amyotrophic lateral sclerosis
(ALS). The current sequence represents the native HK-D3 amino acid

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     comprising the sequence of human kininogen domain 3v defined sequence (S1) of 127 amino acids as given in variant or derivative of native HK-D3 (fully defined
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an isolated anti-angiogenic comprising the sequence of human kininogen domain 3v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tube formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inhibiting angiogenesis, endothelial cell proliferation or endothelial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           79pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specification) or
sequence (S2) of 123
east 20% of activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polypeptide
(HK-D3v) (fi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (fully
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문 δ 밁 Ş 밁 S Matches Query Match Best Local (121 123 123; 61 63 w 2 Similarity EG. AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVDWEKKIYPTVNCQP GKDFVQPTK1CVGCPRD1PTNSPELEETLTHT1TKLNAENNATFYFK1DNVKKARVQVV GKDFVQPPTK1CVGCPRD1PTNSPELEETLTHT1TKLNAENNATFYFK1DNVKKARVQVV Conservative 123 125 98.5%; br 100.0%; Pr Score 660; DB 8; Pred. No. 4.4e-66; Mismatches Length 123; Indels 0; Gaps 122 120 60 62 0

RESULT AD7195 ID AD XX AC AD XX DT 02 XX DE NA XX ON OS HC XX PN US XX PF 1: PA (1) PA ADY71956 standard; protein; 123

ADY71956;

02-JUN-2005

(first entry)

Native human kininogen domain 3 protein.

protein engineering; cancer; cytostatic. kininogen; antiangiogenic; apoptosis; angiogenesis;

Homo sapiens

US2005058599-A1

17-MAR-2005

15-SEP-2003; 2003US-00661784.

15-SEP-2003; 2003US-00661784

(DONA/) DONATE F. (MAZA/) MAZAR A P.

Donate F, Mazar AP

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莱星莱芹芹芹芹菜೩苯위위的的的的的的的的的的的的的数数
                                                                                                                      The invention relates to an isolated anti-antiangiogenic polypeptide (I) can be sequence of human kininogen domain 3 variant (HK-D3v) having a CC fully defined 127 amino acid sequence given in the specification, or its CC fully defined 123 amino acid sequence given in the specification, which CC fully defined 123 amino acid sequence given in the specification, which CC polypeptide, variant or derivative has at least 20% of the activity of CC polypeptide, in inhibiting angiogenesis, endothelial cell proliferation CC or endothelial tube formation in an in vitro or in vivo bioassay. (I) is CC or endothelial tube formation in an in vitro or involves contacting CC useful for inhibiting cell migration, cell invasion, cell proliferation CC cells associated with undesired cell migration, invasion, proliferation CC comprising (I) is useful for treating a subject having a disease or condition associated with undesired cell migration, invasion, condiferation or angiogenesis. (VI) or (VII) is useful for providing to a condition associated with undesired cell migration, invasion, cell; tissue or organ an angiogenesis-inhibitory amount of HK-D3, HK-D3v cell, tissue or organ an angiogenesis-inhibitory amount of HK-D3, HK-D3v corresponds to the native HK-D3 protein sequence.
Query Match
Best Local S
Matches 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel human kininogen domain 3 variant inhibiting angiogenesis, endothelial cell proliferation or endothelial tube formation, useful treating disease associated with undesired cell migration, invasion, proliferation or angiogenesis.
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                                                                                          Sequence 123
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                              Similarity
       98.5%;
ilarity 100.0%;
Conservative (
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                     Score 660; DB 5; pred. No. 4.4e-66;
                                                                    Length 123;
                              Indels
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ADP43683 standard;
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                                                                                       GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
                                                       ĠĸĎĖVQPĖTKI CVĠĊĖRĎI PTŃŚPĖĽEĖTĽTHTI TKLNAĖNNATĖVĖKI ĎNVKKARVQVV
                                            LGM 125
                                LGM 123
  protein;
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RESULT 6
ADP43 SET

ADP45 SET

XX ADP44
XX ADP4
XX ADP4
XX ADP4
XX ADP4
XX Hums
AC Hum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gastrointestinal; cardiovascular; immunosuppressive; antiinflammatory; cytostatic; neuroprotective; gynaecological; gene therapy; gastrointestinal disorder; cardiovascular disorder; autoimmune disorder; inflammatory disorder; cell proliferative disorder; disorder; developmental disorder; epithelial disorder; neurological disorder;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reproductive
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05-DEC-2002;
17-DEC-2002;
24-JAN-2003;
30-JAN-2003;
21-FEB-2003;
                                                                                                                                                                                                                                                                                                                                                      Becha SD,
Kable AE,
                                                                                                                                                                                                                                                                                                New human protein modification and maintenance molecules and polynucleotides for diagnosing, preventing or treating diseases associated with aberrant protein expression, e.g. cardiovascular or cell
                                                                                                                                                                                                                                                                                                                                                                       (INCY-) INCYTE CORP.
                                                                                                                                                                                                                                                                                                                              WPI; 2004-468843/44.
                                                                                                                                                                                                                                                                                                                                          Jin P, Lee S;
                                                                                                                                                                                                                                                                                               polynucleotides for diagnosing, associated with aberrant proteil
                                                                                                                                                                                                                                                                                                                         N-PSDB;
                                                                                                                                                                                                                                                                                                                                                 Burrill
                                                                                                                                                                                                                                                                                          proliferative disorders.
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Marquis
D, Blake
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2003US-0444141P.
2003US-0449491P.
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2002US-0434315P.
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Khare R, Lee SY, Ti
Ho A, Zheng W, Gao
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Gao J, Cha
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                                                                                                                                                                                                                                                                                                                                                                   Richardson TW;
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wla NK,
                                                                                                                                                                                                                                                                                                                                                         Mason PM;
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Claim 1; SEQ ID NO 40; 268pp; English.

commaintenance molecule (PMMM) (II), designated pmodufication and commaintenance molecule (PMMM) (II), designated pmodufication and isolated polynucleotide encoding (I); (2) a recombinant polynucleotide in comprising a plantame of the properties of the polynucleotide in comprising a plantame comprising the recombinant polynucleotide in comprising a plantame comprising the recombinant polynucleotide (I); (3) a cellar corporation with the recombinant polynucleotide (I); (3) a cellar corporation of the recombinant polynucleotide or (I); (3) a cellar corporation of the recombinant polynucleotide or (I); (3) a cellar corporation or purifying (I); (6) an isolated antibody that (I); (6) an isolated antibody that (II); (7) detecting a target polynucleotide; (8) an isolated with decreased expression or treating diseases or conditions associated with decreased expression or treating diseases or condition or an antibody, and an excliptent; (9) that is effective as an agonist or antegonist (II), that specifically that (II); and the specifically of a test compound; (12) a disgnosting for a compound (III) and the specifically of a test compound; (12) a disgnosting test for a condition or disease associated with the expression of promotion antibody with a subject; (II) preparing a polyclonal or disease associated with the expression of promotional antibody with (II); (II); as condition or disease associated with the expression of promotional antibody, and a subject; (II) preparing a polyclonal or golyclonal or monoclonal antibody with the specifically of the antibody in (6); (15) a compositional comprising the polyclonal promotional antibody, and a subject; (II) preparing a polyclonal antibody, and a subject (II) preparing a polyclonal antibody, and a subject (II) and array comprising off a sample containing the compositions of the starget polynucleotial of a sample containing the compositions of the sample containing the compositions of the sample containing the cylonestate, and (IS) an array comprises an intentions associat protein-protein interactions,

268 AA

WO2004053068-A2

24-JUN-2004

03-DEC-2003; 2003WO-US038573

Query Match Best Local Similarity Conservative 98.5%; Score 660; DB 8; 1 100.0%; Pred. No. 1.3e-65; Mismatches Length 268; Indels 0 0

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RESULT 7
ADM83721
ADM9321
A
The invention relates to novel diagnostic and therapeutic polynucleotides CC selected from one of the 2722 sequences defined in the specification. A CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polypoptides may be used to diagnose a particular condition, disease or disorder associated CC with human molecules, e.g. cell proliferative disorder, endocrine CC autoimmune/inflammatory disorder, developmental disorder, endocrine CC disorder, neurological disorders, gastrointestinal disorder, endithp cCC infections caused by virus, bacteria, fungi or parasite. The dithp cCC molecules may also be used in genetic mapping, in identifying individuals CC from minute biological samples, in detecting single nucleotide collymorphisms, as molecular weight markers, and for somatic or germline cCC gene therapy. The present sequence represents a dithp protein of the cCC invention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly
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Lagace RE, Sp. Xu Y, Kwong M, Po
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Stevens KA,
Peralta CH,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Schmidt JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12-SEP-2002; 2002US-0410259P 12-SEP-2002; 2002US-0410260P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human diagnostic and therapeutic pprotein SEQ ID NO:3970
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DB; ACN42373.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INCYTE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GKDFVQPFTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page; 190pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin
Anderson SB, Rioux P, Shen EJ, Work, Stuve LL;
Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton
ng M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, G
Shi X, Suarez CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Wright RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Marjanovic MM,
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Pitte SJ, bic.
Petty TP;
Persti
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Best Local
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15-MAR-2002;
05-APR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24-AUG-2001;
14-DEC-2001;
18-JAN-2002;
                                                                                                                                                                                 New human extracellular messenger (EXMES) polypeptide, useful for preparing a composition for treating a disease associated with decreased expression or overexpression of functional EXMES e.g. autoimmune
                                                                                                                                                                                                                                                                                                                                                                          Ding L, Yao M
Richardson TW,
                                                                                                                                                                  expression or overexpression disorders or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; extracellular messenger; EXMES; cytostatic; antidiabetic; immunosuppressive; autoimmune disorder; inflammatory disorder; diabetes; endocrine disorder; cancer.
                                                                                                                                                                                                                                                                                                                                                      Ramkumar J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003018612-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABP70801;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (INCY-)
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                                                                                                                                                                                                                                                                                                                                                                                                                          SY,
                                                                                                                                                                                                                                                                                  2003-278643/27.
DB; ACC42388.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      63
                                                                                                                                                                                                                                                                                                                                                                                                                                                   BM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INCYTE GENOMICS
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                                                                                                                                                                                                                                                                                                                                                                                                Yao MG,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Lee S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; 2001US-0314811P.
2001US-0340584P.
2002US-0350595P.
2002US-0363432P.
2002US-0363432P.
2002US-036760P.
2002US-0370761P.
2002US-0391378P.
                                                                                                                                                                                                                                                                                                                                                 ee S, Baughn MR, Haler H
re R, Wilson AD, Jin P, H
re R, Welcha SD, Tran UK, C
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Pred. No. 1.3e-65;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Hafalia AJA,
                                                                                                                                                                                                                                                                                                                                                                       , Hawkins PR, Tang YT, B
, Chien D, Zebarjadian Y;
Swarnakar A, Lehr-Mason P
                                                                                                                                                                                                                                                                                                                                                                                                                               Walia NK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 268;
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proteins (EXMES-1 to (ACC42361-ACC42388).

Page

224pp;

English

The present invention relates to novel human extracellular messenger proteins (EXMES-1 to-28; ABP70774-ABP70801) and their coding sequence (ACC42361-ACC42388). The proteins are useful for preparing a composit for diagnosing or treating a disease or condition associated with

sequences composition

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                    Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F;
Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV;
Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH;
Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES
Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gietz
Patury S, Shi X, Suarez CJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-NOV-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
The invention relates to novel diagnostic and therapeutic polynucleotides selected from one of the 2722 sequences defined in the specification. A polynucleotide of the invention may have a use in gene therapy. The human diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated with human molecules, e.g. cell proliferative disorders,
                                                                                                                                                                                                                              New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or in gene mapping.
                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-329368/30.
N-PSDB; ACN42372.
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12-SEP-2002; 2002US-0410260P.
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                                                                                                                                                                                      Claim 27;
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                                                                                                                                                                                      Page; 190pp;
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                                                                                                                                                                                        English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Stuve LL;
UA, Kirton ES;
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RESULT 10
ABP70799
ID ABP70
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          autoimmune/inflammatory disorder, developmental disorder, endocrine disorder, neurological disorders, gastrointestinal disorders, or infections caused by virus, bacteria, fungi or parasite. The dithp molecules may also be used in genetic mapping, in identifying individuals from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline gene therapy. The present sequence represents a dithp protein of the invention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm
                                                                                                                                                                    24-AUG-2001; 2001US-0314811P.
14-DEC-2001; 2001US-0340584P.
18-JAN-2002; 2002US-0350595P.
11-MAR-2002; 2002US-0363432P.
15-MAR-2002; 2002US-0364607P.
05-APR-2002; 2002US-0370761P.
24-JUN-2002; 2002US-0391378P.
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                                                                                                                                                                                                                                                                                                                                                                                Human; extracellular messenger; EXMES; cytostatic; antidiabetic; immunosuppressive; autoimmune disorder; inflammatory disorder; c
                                                                                                                                                                                                                                                                                                                                                                                                                     Human extracellular messenger,
                                                                                                                                                                                                                                                                                                                                                                                                                                              26-AUG-2003
                                                                                                                                                                                                                                                                                              06-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                      endocrine disorder; cancer.
                                                                                                                                                                                                                                                                     22-AUG-2002; 2002WO-US027213
                                                                                                                                                                                                                                                                                                                     WO2003018612-A2
                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                    Ding L, Yao M
Richardson TW,
                                                                                                                                                                                                                                                                                                                                                                               immunosuppressive; autoimmune
                                   N-PSDB;
                                                                                                                                                (INCY-)
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                                   2003-278643/27.
)B; ACC42386.
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                                                                                             Khare R,
Yao MG,
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ilarity 100.0%;
Conservative
                                                                                    Lee S, Baughn MR, Hafalia AJA, We
hare R, Wilson AD, Jin P, Hawkins I
ao MG, Becha SD, Tran UK, Chien D,
TW, Kable AE, Chang H, Swarnakar J
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Pred. No. 1.5e-65;
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                                                                                       , Walia NK, Elliou ...
ins PR, Tang YT, Burfc
in D, Zebarjadian Y;
Thr-Mason PM;
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                                                                                                                              Elliott VS;
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                                                                                                                 Burford
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New human extracellular preparing a composition

messenger (EXMES) polypeptide, useful for treating a disease associated with

with

decreased

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RESULT 11
ABM83717
ID ABM83717
ID ABM83717
ID ABM83717
ID ABM83717
AC ABM83717
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Best Local S
Matches 123
                                                                                                                                                                                                                                                                    Harthshorne Mooney EM, I
Stevens KA,
Peralta CH,
New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy of in gene mapping.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-SEP-2002;
12-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                 Schmidt JP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human diagnostic and therapeutic pprotein SEQ ID NO:3966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-NOV-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 322 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to novel human extracellular messenger proteins (EXMES-1 to-28; ABP70774-ABP70801) and their coding sequences (ACC42361-ACC42388). The proteins are useful for preparing a composition for diagnosing or treating a disease or condition associated with decreased expression or overexpression of functional EXMES e.g. autoimmune/inflammatory disorders, diabetes, endocrine disorders or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              expression or overexpression of functional EXMES e.g. autoimmune disorders or cancer.
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123; Conser
                                                                                                                                                                                        It JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F; horne TA, Suchorolski MT, Altus CM, Pitte SJ, Elder LV EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP; s KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin a CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL; RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, GS, Shi X, Suarez CJ;
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2002US-0410260P
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Best Local Similarity
                                               12-OCT-2001;
25-OCT-2001;
09-NOV-2001;
28-NOV-2001;
06-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cell proliferation disorder; atherosclerosis; neurological disorder; epileps; Huntington's disease; stroke; immune disorder; allergy; inflammatory disorder; AIDS; developmental disorder; hypothyroidism; cushing's syndrome; gastrointestinal disorder; epithelial disorder; infection; cytostatic; antiarteriosclerocic; anticonvulsant; nootropinfection; cytostatic; anticonvulsant; nootropinfection; no cytostatic; no cytostatic; no cytostatic; no cyto
                                                                                                                                                                                                                         11-OCT-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; protein modification and maintenance molecule;
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                    ; 2001US-0329689P.
; 2001US-0335703P.
; 2001US-0334887P.
; 2001US-0334145P.
; 2001US-0337451P.
; 2001US-0337454P.
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ilarity 100.0%;
Conservative
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The present invention relates to the isolation of human protein CC modification and maintenance molecules (PMAM), and the polymucleotide CC sequences encoding them. A total of 40 PMAM polypeptides (designated PMAM CC -1 to PMAM-40) are disclosed. The sequences of the invention are useful CC for diagnosing a condition or disease associated with the expression of CC for diagnosing a condition or disease associated with the expression of CC pMAM in a subject, preparing a polyclonal or monoclonal antibody, and CC generating an expression profile of a sample containing the CC expression or overexpression of PMAM are cell proliferation disorders (e.g. epilepsy, CC (e.g. cancer, atherosclerosis), neurological disorders (e.g. epilepsy, CC (e.g. cancer, atherosclerosis), neurological disorders, (e.g. epilepsy, CC allergies), developmental disorders (e.g. hypothyroidism, Cushing's syndrome), gastrointestinal or epithelial disorders, and infections. The EpMAM polypeptides or their fragments are useful in screening compounds that compounds the target polymucleotide and compounds that grecifically bind to, or modulate the activity of the polypeptide. ABU9201-ABU92060 represent the human PMAM polypeptides of the invention of the target polymucleotide in the invention of the 
                                                                                                                                                                                                                                                                                                                RESULT 13
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Best Local S
Matches 123
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Tran UK, Becha SD, Duggan BM, Lee EA, Griffin JA, Li JX;
Sprague WW, Hafalia AJA, Chawla NK, Lehr-Mason PM, Kable AE,
Marquis JP, Yao MG, Richardson TW, Tang TY, Jin P, Chien D;
Bhatia U, Burrill JD, Lee S, Blake JJ, Ho A, Zheng W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New human protein modification and maintenance molecules (PMMM), useful for diagnosing, treating and preventing diseases or conditions associat with the aberrant PMMM expression e.g. cancer, atherosclerosis, or
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N-PSDB; ACA92439.
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The invention relates to novel diagnostic and therapeutic polynucleotides comply the control of the 2722 sequences defined in the specification. A comply control of the invention may have a use in gene therapy. The human comply control of the condition of the properties may be comply to diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be comply to the condition, disease or disorder associated comply the condition, disease or disorder associated comply the condition of the condition of the condition of the conditions caused by virus, bacteria, fungion parasite. The dithp completes may also be used in genetic mapping, in identifying individuals comply comply as molecular weight markers, and for somatic or germline comply comply conditions. The present sequence represents a dithp protein of the comparition. Note: The sequence represents a dithp protein of the comparition. Note: The sequence comply com
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12-SEP-2002; 2002US-0410260P.
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                      Human extracellular messenger, EXMES-27
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                                                                                                                                                                     standard; protein;
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Suchorolski MT, Altus CM, Pitts SJ, Blde
Suchorolski MT, Altus CM, Pitts SC, Reddy
                                                                            entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1.7e-65;
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14-DEC-2001; 2001US-0340584P.
18-JAN-2002; 2002US-0350595P.
11-MAR-2002; 2002US-0363432P.
15-MAR-2002; 2002US-0364607P.
05-APR-2002; 2002US-0370761P.
24-JUN-2002; 2002US-0391378P.
             08-SEP-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 preparing a composition for treating a disease associated with de expression or overexpression of functional EXMES e.g. autoimmune disorders or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                      The
                                      AEB32339
                                                              AEB32339 standard;
                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                     for diagnosing or treating a disease or condition associated wi decreased expression or overexpression of functional EXMES e.g. autoimmune/inflammatory disorders, diabetes, endocrine disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 206; 224pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to novel human extracellular messenger proteins (EXMES-1 to-28; ABP70774-ABP70801) and their coding sequences (ACC42361-ACC42388). The proteins are useful for preparing a composition for diagnosing or treating a disease or condition associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (INCY-)
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SY, Khare R, Wilson AD, Jin P, Hawkins PR, Tang YT, Bux
g L, Yao MG, Becha SD, Tran UK, Chien D, Zebarjadian Y,
hardson TW, Kable AE, Chang H, Swarnakar A, Lehr-Mason PM,
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DB; ACC42387.
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100.0%; Pred. No. 1.9e-65;
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AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP

307

GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV

GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV

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Query Match Best Local S Matches 123

Similarity

98.5%;

Score 660; DB 9; Pred. No. 1.9e-65;

Length 362; Indels

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Conservative

<u>.</u>.

Mismatches

Sequence

362

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The invention relates to a detection reagent capable of detecting one or composed more single nucleic acid polymorphisms. The invention also relates to concentration whether a trait is linked too one of the human chromosomes or composed methor a trait is linked too one of the human chromosomes or composed information given in the specification, an isolated nucleic acid molecule for detecting at least one SNP given in the specification comprising at least about 12 contiguous nucleotides, gencyping at least one containing the detection reagent. CC individual having or at risk of developing a disorder and a kit comprising at least one containing whether a trait is linked to one of the human chromosomes or its sub-region comprises determining whether the trait is linked to one comprises contacting the detection reagents. Genotyping at least one SNP position given in the specification in a sample comprises contacting the sample with a detection reagent that differentiates between alternative comprises genotyping at least one SNP position given in the specification. CC alleles at at least one SNP position given in the specification in a comprise segmentyping at least one SNP position. CC identifying an individual having or at risk of developing a disorder comprise segmentyping at least one SNP given in the specification in a comprise segment promote the individual. The disorder is type II diabetes consider in the specification reagent is considered at for this patent diabetes or obesity. This sequence considered at for this patent diabetes or obesity. This sequence considered and cc specification but was obtained in electronic format from USPTO at the sequence and cc sequence hatmines and considered in electronic format from USPTO at the considered at the sequence had not form part of the printed at the considered and conside
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 200
N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New detection reagent capable of detecting 1, 100, 500, 1000 or 5000 more single nucleic acid polymorphisms, useful in identifying an individual having or at risk of developing type II diabetes or obesi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 102; 31pp; English.
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10-SEP-2001; 2001US-00948947.
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ALIGNMENTS

RESULT 2 US-09-949-002-476 ; Sequence 476, A ; Patent No. 6900 ; GENERAL INFORMA ; APPLICANT: VENE ; TITLE OF INVEN ; TITLE OF INVEN ; TITLE OF INVEN	Qy Db	D Q	₽ &	Query Match Best Local Matches 12	RESULT 1 US-09-461-061-18 J Sequence 18, A J Sequence 18
-09-949-002-476 -09-949-002-476 Sequence 476, Application US/09949002 Sequence 400016 GENERAL INFORMATION: APPLICANT: VENTER, J. Craig et al. TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED TITLE OF INVENTION: WITH INFLAMMATORY AUTOIMMUNE DISEASE, METHODS OF DETECTION TITLE OF INVENTION: AND USES THEREOF	123 LGM 125 121 LGM 123	63 AGKKYPIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP 122	3 GKDFVQPFTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV 62 	<pre>/ Match 98.5%; Score 660; DB 2; Length 123; Local Similarity 100.0%; Pred. No. 1e-65; les 123; Conservative 0; Mismatches 0; Indels 0; Gaps 0;</pre>	SULT 1 -09-461-061-18 Sequence 18, Application US/09461061 Patent No. 6889931 GENERAL INFORMATION: Inhibition of Angiogenesis By High Molecular Weight TITLE OF INVENTION: Inhibition of Angiogenesis By High Molecular Weight TITLE OF INVENTION: Kininogen Domain 3 Peptide Analogs FILE REPERENCE: 6056-260 US CURRENT APPLICATION NUMBER: US/09/461,061 CURRENT FILING DATE: 1999-12-15 EARLIER APPLICATION NUMBER: 60/112,427 EARLIER FILING DATE: 1998-12-16 NUMBER OF SEQ ID NOS: 21 SOPTWARE: Patentin Ver. 2.0 SEQ ID NO 18 LENGTH: 123 TYPE: PRT ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Description of Artificial Sequence: Human HK OTHER INFORMATION: domain 3

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RESULT 4
US-09-919-039-29
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US-09-949-002-480
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; Sequence 29, Application US/09919039
; Patent No. 6727066
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US-09-949-002-476
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH INFLAMMATORY AUTOIMMUNE DISBASE, METHODS OF DETECTION
TITLE OF INVENTION: AND USES THEREOF
FILE REPERENCE: CLOO0790
CURRENT APPLICATION NUMBER: US/09/949,002
CURRENT FILING DATE: 2000-01-28
PRIOR FILING DATE: 2000-09-08
RIUMBER OF SEQ ID NOS: 10823
SOFTWARE: FASESEQ for Windows Version 4.0
SEQ ID NO 480
LENGTH: 362
TYPE: PRT
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LENGTH: 362
TYPE: PRT
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Best Local Similarity 100.0%; Pred. No. 4.3e-65;
Matches 123; Conservative 0; Mismatches 0;
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Best Local &
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CURRENT APPLICATION NUMBER: US/09/949,002
CURRENT FILING DATE: 2000-01-28
PRIOR APPLICATION NUMBER: 60/231,401
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 10823
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mes 123; Conserv
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100.0%; Pred. No. 4.3e-65;
Live 0; Mismatches 0;
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; IENGTH: 427
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc feature
; OTHER INFORMATION: Incyte II
US-09-919-039-29
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FIITLE OF INVENTION: GENES EXPRESSED IN TREATED HUN-
FILE REFERENCE: PA-0035 US

CURRENT APPLICATION NUMBER: US/09/919,039

CURRENT FILING DATE: 2002-09-09

PRIOR APPLICATION UNMBER: 60/222,113

PRIOR FILING DATE: 2000-07-28

NUMBER OF SEQ ID NOS: 401

SOPTWARE: PERL Program

SEQ ID NO 29

SEQ ID NO 29
                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/949,002
CURRENT FILING DATE: 2000-01-28
PRIOR APPLICATION NUMBER: 60/231,401
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 10823
SOFTWARE: FASUSEQ for Windows Version 4.0
SEQ ID NO 475
LENGTH: 579
TYPE: PRT
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US-09-949-002-475
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US-09-949-002-475
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APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH INFLAMMATORY AUTOIMMUNE DISEASE, METHODS OF DETECTION

TITLE OF INVENTION: AND USES THEREOF

FILE REFERENCE: CL000790
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Best Local Similarity
Matches 123; Conserv
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Best Local Similarity
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Patent No. 6900016
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       LGM 125
                                         AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
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Pred. No. 8e-65;
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308

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US-08-193-114B-1
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APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH INFLAMMATORY AUTOIMMUNE DISEASE, MITTILE OF INVENTION: AND USES THEREOF
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SOFTWARE: FRETSEQ for Windows Version 4.0
SEQ ID NO 481
LENGTH: 579
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Best Local 9
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PRIOR APPLICATION NUMBER: 60/231,401
PRIOR FILING DATE: 2000-09-08
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CURRENT APPLICATION NUMBER: US/09/949,002
CURRENT FILING DATE: 2000-01-28
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APPLICANT: Jiang, Yongping
TITLE OF INVENTION: Modulation of Blood
TITLE OF INVENTION: Pressure and Inhibition of Platelet Activation
TITLE OF INVENTION: with Kininogen Fragment
NUMBER OF SEQUENCES: 2
                                                         APPLICATION NUMBER: US/08/193,114B
FILING DATE: 9 February 1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 720
COMPUTER: IBM PS/2
                                                                                                                                                  SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                       STREET: 1800 Two Penn Center Plaza
CITY: Philadelphia
STATE: Pennsylvania
COUNTRY: U.S.A.
                                                                                                                                                                                              OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                  ZIP: 19102
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             APPLICATION NUMBER: U.S. Application
APPLICATION NUMBER: Serial No. 5472945 07/744,545
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13 August 1991
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100.0%; Pr
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PCT-US92-06809-1
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                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch, 720 Kb
COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WordPerfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US92/06809
FILING DATE: 19910813
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Best Local Similarity
                                                                                                                                       CLASSIFICATION: 514

PRIOR APPLICATION DATA:
APPLICATION NUMBER: U.S. Application
APPLICATION NUMBER: Serial No. 744,545

FILING DATE: 13 August 1991
ATTORNEY/AGENT INFORMATION:
NAME: MONACO, DANIEL A.
REGISTRATION NUMBER: 30,480
REFERENCE/DOCKET NUMBER: 6056-137
TELEPHONE: (215) 568-8383
TELEPHONE: (215) 568-5549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IELEX: NO. 5472945e.
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 117 amino acids
TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: MONACO, Daniel A.
REGISTRATION NUMBER: 30,480
REFERENCE/DOCKET NUMBER: 6056
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-8383
TELEFAX: (215) 568-5549
                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 117 amino acid
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Schmaier, Alvin H. APPLICANT: Jiang, Yongping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Temple University - Of the ADDRESSEE: Commonwealth System of Higher Education STREET: 406 University Services STREET: Building CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: peptide TOPOLOGY: line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14 CVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKYFIDFVA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 CVGCPRDIPTNSPBLEBTLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKYFIDFVA 60
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': U.S.A.
                                                         117 amino acids
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: Modulation of Blood

: Pressure by Altering Bradykinin Levels
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100.0%; Pred. No. 4.5e-59;
htive 0; Mismatches 0;
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Best Local Similarity
                                                                                                                                                    Query Match
Best Local 9
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                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                       TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US
FILLING DATE: Filed Her
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: NOVEL HUMAN CYSTATIN-LIKE TITLE OF INVENTION: PROTEIN
                                                                                                                                                                                                                                     IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: DOS
SOFTWARE: FABISEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
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CITY: Palo Alto
                                                                                                                                                                                                                                                                LENGTH: 178 amino
TYPE: amino acid
STRANDEDNESS: sir
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                                                                                                                                                  Local
                                                                                                                                                                                                                       CLONE: 30443
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                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE:
114 VEIGRTTCKKNQHLRL-DDCDFQTNHTLKQTLSCYSEVWVVPW----VPALRGACSPLSL 168
                                                                   54
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                                                                                               11 TKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKYFID 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94304
                                                                                                                                 h 25.2%; Score 169; DB 1; Length 178; Similarity 32.5%; Pred. No. 7.2e-11; 39; Conservative 22; Mismatches 49; Indels
                 FVARETTCSKESNEELTESCE---TKKLGQSLDCNAEVYVVPWEKKIYPTVN---CQPLGM 125
                                                               SRVKPGFPKTIKTNDPGVLQAARYSVEKFNNCTNDMFLFKESRITRALVQIVKGLKYMLE 113
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                                                                                                                                    10;
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                                                                                                                                                                                                                                 US-09-461-061-17
                                                                                                                                                                                                                                                    RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-314-777-1
Sequence 17, Application US/09461061
Patent NO. 6869931
GENERAL INFORMATION:
APPLICANT: McCrae, Keith R.
TITLE OF INVENTION: Inhibition of Angiogenesis By High Molecular Weight TITLE OF INVENTION: Kininogen Domain 3 Peptide Analogs
FILE REFERENCE: 6056-260 US
CURRENT APPLICATION NUMBER: US/09/461,061
CURRENT FILING DATE: 1999-12-15
EARLIER APPLICATION NUMBER: 60/112,427
EARLIER FILING DATE: 1998-12-16
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OPERATING SYSTEM: DOS
SOTTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
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APPLICANT: Goli, Surya K.
TITLE OF INVENTION: NOVEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS LENGTH: 178 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER:
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CORRESPONDENCE ADDRESS:
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CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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                                                                                                                                                                                                                                                                                                               114 VEIGRTTCKKNOHLRL-DDCDFQTNHTLKQTLSCYSEVWVVPW----VPALRGACSPLSL
                                                                                                                                                                                                                                                                                                                                                                                              54 SRVKPGFPKTIKTNDPGVLQAARYSVEKFNNCTNDMFLFKESRITRALVQIVKGLKYMLE 113
                                                                                                                                                                                                                                                                                                                                                                                                                                      11 TKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKYFID 70
                                                                                                                                                                                                                                                                                                                                                 71 FVARETTCSKESNEELTESCE---TKKLGQSLDCNAEVYVVPWEKKIYPTVN--CQPLGM 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ch 25.2%; Sc
l Similarity 32.5%; Pr
39; Conservative 22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 169; DB 2;
Pred. No. 7.2e-11;
2; Mismatches 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 178;
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NUMBER OF SEQ ID NOS:

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                                                                                                                              Best Loca
Matches
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Patent No. 5919658
GENERAL INFORMATION:
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SEQ ID NO 17
LENGTH: 32
                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: Analog OTHER INFORMATION: human HK domain 3 fragment
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS LENGTH: 145 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA; APPLICATION NUMBER: US/08/832,535 FILING DATE: 03-APR-1997 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0
                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: (201) 994-1700
                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: KIMBALL, PAUL C.
REGISTRATION NUMBER: 34,6
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                           TYPE: amino acid
TOPOLOGY: linear
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mes 31; Conserv
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                              71 FVARETTCSKESNEELTESCE---TKKLGQSLDCNAEVYVVPWEKKI-YPTVNC 120
                                                                                                11 TKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKYFID 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 87 TESCETKKLGOSLDCNAEVYVVPWEKKIYPTV 118
92 VEIGRTTCKKNOHLRL-DDCDFQTNHTLKQTLSCYSEVWVVPWLQHFEVPVLRC 144
                                                                 SRVKPGFPKTIKTNDPGVLQAARYSVEKFNNCTNDMFLFKESRITRALVQIVKGLKYMLE
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                                                                                                                                                                                                                                                               145 amino acids
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Pred. No. 1.6e-11;
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GENERAL INFORMATION:

91

APPLICANT:

Li, Haodong Yu, Guo-Liang

Gentz, Reiner

APPLICANT:

APPLICANT: Ni, Jian
TITLE OF INVENTION: Cystatin
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:

CITY: Rockville

9410 Key West Avenue

Human Genome Sciences,

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ADDRESSEE:

COUNTRY:

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US-09-019-485-2
                                   US-09-019-485-3
Patent No.
              Sequence 3,
                                                                                                                                                                                                                                                       Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Li, Ha
APPLICANT: Yu, Gu
                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 3013098439
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Cystatin F NUMBER OF SEQUENCES: 17 CORRESONE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Benson, Robert H.
REGISTRATION NUMBER: 30,446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
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                                                                                                        92 VEIGRTTCKKNOHLRL-DDCDFQTNHTLKQTLSCYSEVWVVPWLQHFEVPVLRC 144
                                                                                                                                      71 FVARETTCSKESNEELTESCE---TKKLGQSLDCNAEVYVVPWEKKI-YPTVNC 120
                                                                                                                                                                              32 SRVKPGFPKTIKTNDPGVLQAARYSVEKFNNCTNDMFLFKESRITRALVQIVKGLKYMLE
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                Application US/09019485
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9410 Key West Avenue
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Ni, Jian
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3013098504
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ZIP: 20850 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk

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APPLICANT: Hollows, James L.

APPLICANT: Feldhaus, Andrew

TITLE OF INVENTION: TESTIS SPECIFIC CYSTATIN-LIKE PROTEIN CYSTATIN T

FILE REFERENCE: 95-72

CURRENT APPLICATION NUMBER: US/09/431,480

CURRENT FILING DATE: 1999-11-01

EARLIER APPLICATION NUMBER: 60/109,217

EARLIER APPLICATION NUMBER: 60/156,382

EARLIER APPLICATION NUMBER: 60/156,382

EARLIER FILING DATE: 1999-09-28

NUMBER OF SEQ ID NOS: 22

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 9

LENGTH: 145

TYPE: PRT

ORGANISM: Homo sapiens

US-09-431-480-9
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; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-09-019-485-3
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                                                                                                                                                                                    Query Match 24.7%; Score 165.5; DB 2; Length 145; Best Local Similarity 32.5%; Pred. No. 1.3e-10; Matches 37; Conservative 22; Mismatches 50; Indels 5.
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INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Benson, Robert H.
REGISTRATION NUMBER: 30,446
REFERENCE/DOCKET NUMBER: PF265P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 3013098504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/019,485
FILING DATE:
CLASSIFICATION:
92 VEIGRTTCKKNOHLRL-DDCDFQTNHTLKQTLSCYSEVWVVPWLQHFEVPVLRC 144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   71 FVARETTCSKESNEELTESCE---TKKLGQSLDCNAEVYVVPWEKKI-YPTVNC 120
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Search completed: February 8, 2006, 03:18:49
Job time : 48 secs

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Result
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length: 2000000000
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291.780 Million cell updates/sec
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Maximum Match 100%
Listing first 45 summaries
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
   GenCore version (c) 1993 - 2006
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US-10-893-315-102
US-10-637-313-20
US-10-637-313-24
US-10-637-313-24
US-10-637-313-24
US-10-637-313-24
US-10-893-315-105
US-10-893-315-105
US-10-893-315-105
US-10-637-313-14
US-10-637-313-14
US-10-637-313-18
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102, App
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20, Appl
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24, Appl
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105, Appl
106, Appl
107, Appl
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5 US-10-637-313-56 5 US-10-637-313-58 5 US-10-637-313-62 5 US-10-637-313-62 5 US-10-637-313-64 5 US-10-637-313-66 5 US-10-41-600-1180 5 US-10-450-763-51460 5 US-10-637-313-72 5 US-10-637-313-72 5 US-10-637-313-74 5 US-10-637-313-74 5 US-10-637-313-74 5 US-10-637-313-74 5 US-10-637-313-74 5 US-10-637-313-74	357	357	435	127	644	644	644	644	644	644	644	644	644	644	644	644	644	611
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	US-10-637-313-46	US-10-637-313-10	US-10-450-763-51464	US-10-661-784-3	US-10-637-313-74	US-10-637-313-72	US-10-637-313-70	US-10-637-313-68	US-10-637-313-66	US-10-450-763-51460	US-10-741-600-1180	US-10-637-313-76	US-10-637-313-64	US-10-637-313-62	US-10-637-313-60	US-10-637-313-58	-10-637-313	CC +C CL C+C C+
	46, App	10, Appl	51464,	3, Appl	74, App	72, App	•	68, Appl	•	51460,	1180, A	76, App	64, App	62, Appl	60, App	58, App	56, App	A. 1. 1. C.

ALIGNMENTS

RESULT 1 US-10-661-784-2

Sequence 2, Application US/10661784 Publication No. US20050058599A1 GENERAL INFORMATION:

APPLICANT: DONATE, Fernando APPLICANT: MAZAR, Andrew

TITLE OF INVENTION: Human Kininogen D3 Domain Polypeptide as an Anti-Angiogenic and TITLE OF INVENTION: Anti-Tumor Agent FILE REFERENCE: 38942-193024
CURRENT APPLICATION NUMBER: US/10/661.784
CURRENT FILING DATE: 2003-09-15
PRIOR APPLICATION NUMBER: US 60/401,279
PRIOR APPLICATION NUMBER: US 60/401,279
PRIOR PILING DATE: 2002-09-13
PRIOR PILING DATE: 2002-09-13
PRIOR PILING DATE: 2002-09-13

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RESULT 2
US-10-661-784-1
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           Sequence 1, Application US/10661784
Publication No. US20050058599A1
GENERAL INFORMATION:
APPLICANT: DONATE, Fernando
APPLICANT: MAZAR, Andrew
TITLE OF INVENTION: Human Kininogen D3 Domain
TITLE OF INVENTION: Anti-Tumor Agent
FILE REFERENCE: 38342-193024
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Matches
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CURRENT APPLICATION NUMBER: US/10/661,784
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                                                                                                                                                                                                                                                                                                                                                                                                        1 GSGKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQ
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                                                     Polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 125;
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                                                     Anti-Angiogenic
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FILE REFERENCE: CL000786
CURRENT APPLICATION NUMBER: US/10/893,315
CURRENT FILING DATE: 2004-07-19
PRIOR APPLICATION NUMBER: 60/231,397
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 2172
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 102
LENGTH: 362
TYPE: PAT
ORGANISM: Human
US-10-893-315-102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: US 60/
PRIOR FILING DATE: 2002-09-13
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin version 3.2
SEQ ID NO 1
LENGTH: 123
TYPE: PRT
ORGANISM: Homo sapiens
             RESULT 4
US-10-893-315-104
; Sequence 104, Application US/10893315
; Publication No. US20050147987A1
; GENERAL INFORMATION:
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US-10-893-315-102
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Publication No. US20050147987A1

GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN

TITLE OF INVENTION: WITH TYPE II DIABETES

TITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
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Best Local Similarity
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VENTER, J. Craig et al
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                                                                                                                                                                                                                                                                                                                   98.5%; Score 660; DB 5; 100.0%; Pred. No. 2.9e-61; tive 0; Mismatches 0;
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; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; TITLE OF INVENTION: WITH TYPE II DIABETES AND OBESITY, METHODS OF DETECTION AND
; TITLE OF INVENTION: USES THEREOF
; FILE REFERENCE: CLOOO736
; CURRENT APPLICATION NUMBER: US/10/893,315
; CURRENT FILING DATE: 2004-07-19
; PRIOR APPLICATION NUMBER: 60/231,397
; PRIOR APPLICATION NUMBER: 60/231,397
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOM: 2172
; SOFTWARE: FASTESEQ for Windows Version 4.0
; SEQ ID NO 104
; LENGTH: 362
; TYPE: PRT
; ORGANISM: Human
US-10-893-315-104
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APPLICANT: Zerhusen, Bryan D.
TITLE OF INVENTION: Therapeutic Polypeptides, Nu.
FILE REFERENCE: 21402-377 B
CURRENT APPLICATION NUMBER: US/10/162,335
CURRENT FILING DATE: 2002-10-01
PRIOR APPLICATION NUMBER: 60/295,607
PRIOR FILING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: 60/295,661
PRIOR PILING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: 60/296,404
PRIOR APPLICATION NUMBER: 60/296,404
PRIOR APPLICATION NUMBER: 60/296,404
PRIOR FILING DATE: 2001-06-06
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US-10-162-335-82
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Best Local Similarity
Matches 123; Conserv
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APPLICANT:
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Casman, Stacie J.
Edinger, Shlomit R.
Gangolli, Esha A.
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Spytek, Kimberly A.
Vernet, Corine A. M.
Voss, Edward Z.
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Pena, Carol E. A.
Rastelli, Luca
Shimkets, Richard A.
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t, Tord
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Valerie
                                                                                                                                                                                                Nucleic Acids Encoding Same, and Method
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 201
SEQ ID NO 82
LENGTH: 390
TYPE: PRT
ORGANISM: Homo sapiens
US-10-162-335-82
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US-10-637-313-32
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CURRENT APPLICATION NUMBER: US/10/637,313
CURRENT FILING DATE: 2003-08-08
PRIOR APPLICATION NUMBER: 10/16235
PRIOR FILING DATE: 2002-06-03
PRIOR FILING DATE: 2002-06-04
PRIOR PPLICATION NUMBER: 60/295607
PRIOR PPLICATION NUMBER: 60/295661
PRIOR PPLICATION NUMBER: 60/296404
PRIOR PILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: 60/296418
PRIOR APPLICATION NUMBER: 60/296418
PRIOR APPLICATION NUMBER: 60/296418
PRIOR PRILING DATE: 2001-06-06
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Best Local Similarity 100.0%; Pred. No. 3.1e-61;
Matches 123; Conservative 0; Mismatches 0;
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APPLICANT:
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APPLICANT:
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APPLICANT: Uriel
APPLICANT: Li Li
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APPLICANT: Charles Miller
APPLICANT: Nikolai Khramtsov
APPLICANT: Corine Vernet
APPLICANT: Corine Vernet
APPLICANT: Meera Patturajan
APPLICANT: M
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OR FILING DATE: 2001-06-12
OR APPLICATION NUMBER: 60/298,285
OR FILING DATE: 2001-06-14
OR APPLICATION NUMBER: 60/298,556
OR FILING DATE: 2001-06-15
OR APPLICATION NUMBER: 60/299,949
OR APPLICATION NUMBER: 60/290,949
OR APPLICATION NUMBER: 60/290,949
OR APPLICATION NUMBER: 60/300,883
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FILING DATE: 2001-06-06
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No. US20040229779A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sasha
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APPLICANT: Zerhusen, Bryan D.
TITLE OF INVENTION: Therapeutic Polypeptides, Nu.
FILE REFERENCE: 21402-377 B
CURRENT APPLICATION NUMBER: US/10/162,335
CURRENT FILING DATE: 2002-10-01
PRIOR APPLICATION NUMBER: 60/295,607
PRIOR APPLICATION NUMBER: 60/295,661
PRIOR APPLICATION NUMBER: 60/296,404
PRIOR APPLICATION NUMBER: 60/296,404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Anderson, David W.
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LENGTH: 390
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Best Local Similarity
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PRIOR FILING DATE: 2001-06-28
Remaining Prior Application data removed
NUMBER OF 8BQ ID NOS: 255
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PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/299949
PRIOR FILING DATE: 2001-06-21
PRIOR APPLICATION NUMBER: 60/300883
PRIOR FILING DATE: 2001-06-26
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                                                                                                                                                                          Stone, David J.
Spytek, Kimberly A.
Vernet, Corine A. M.
Voss, Edward Z.
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                                                                                                                                                                                                                                              Rastelli, Luca
Shimkets, Richard A.
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                                                                                                                                                                                                                                                                                   Patturajan,
Pena, Carol
                                                                                                                                                                                                                                                                                                                                       MacDougall, John
Malyankar, Uriel
Millet, Isabelle
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Hjalt, Tord
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А.
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    See File Wrapper or PALM

                                                                                                                                             Nucleic
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; ORGANISM: Homo
US-10-162-335-70
            APPLICANT: Corine Vernet
APPLICANT: Meera Patturajan
TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUMERICE REFERENCE: 21402-593B
CURRENT APPLICATION NUMBER: US/10/637,313
CURRENT FILING DATE: 2003-08-08
PRIOR APPLICATION NUMBER: 10/162335
PRIOR FILING DATE: 2002-06-03
PRIOR APPLICATION NUMBER: 60/295607
PRIOR FILING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: 60/295661
PRIOR FILING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: 60/2956404
PRIOR APPLICATION NUMBER: 60/296404
PRIOR APPLICATION NUMBER: 60/296418
PRIOR APPLICATION NUMBER: 60/296418
PRIOR FILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: 60/296418
PRIOR FILING DATE: 2001-06-06
PRIOR FILING DATE: 2001-06-06
PRIOR FILING DATE: 2001-06-06
PRIOR FILING DATE: 2001-06-06
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US-10-637-313-20
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Best Local Similarity
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 201
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APPLICANT: Uriel Malyankar
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RR FILING DATE: 2001-06-14
DR APPLICATION NUMBER: 60/298,556
RR FILING DATE: 2001-06-15
DR APPLICATION NUMBER: 60/299,949
RR FILING DATE: 2001-06-21
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RAPPLICATION NUMBER: 60/296,418

RR FILING DATE: 2001-06-06

PR APPLICATION NUMBER: 60/297,414

RAPPLICATION NUMBER: 60/297,567

RAPPLICATION NUMBER: 60/297,567

RAPPLICATION NUMBER: 60/297,567
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APPLICATION NUMBER: 60/300,883
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                                                                                                                                                                                                                                                                                    Stacie Casman
Ferenc Boldog
Charles Miller
Nikolai Khramtsov
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                                                                                                                                                                                                                                                                                                                                                                         Sasha Guo
Mei Zhong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LGM 346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/10637313
No. US20040229779A1
                                                                                                                                                                                                                                                                                                                                                         Muralidhara Padigaru
                                                                                                                                                                                                                                                                                                                                                                                                          David Anderson
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373
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                                 LGM 125
LGM 375
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; SEQ ID NO 20
; LENGTH: 398
; TYPE: PRT
; ORCANISM: Homo sapiens
US-10-637-313-20
                                                                                                                                                                                                                                                                                                   ; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID
US-09-919-039-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: 60/29856
PRIOR APPLICATION NUMBER: 60/299949
PRIOR APPLICATION NUMBER: 60/299949
PRIOR APPLICATION NUMBER: 60/30083
PRIOR FILING DATE: 2001-06-26
PRIOR PRIOR APPLICATION NUMBER: 60/301550
PRIOR APPLICATION NUMBER: 60/301550
PRIOR APPLICATION NUMBER: 60/301550
PRIOR APPLICATION NUMBER: 60/301550
PRIOR FILING DATE: 2001-06-28
PRIOR PRIOR APPLICATION NUMBER: 60/301550
PRIOR PRIOR APPLICATION NUMBER: 60/301550
PRIOR PRIOR DATE: 2001-06-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/919,039
CURRENT FILING DATE: 2002-09-09
PRIOR APPLICATION NUMBER: 60/222,113
PRIOR FILING DATE: 2000-07-28
NUMBER OF SEQ ID NOS: 401
SOFTWARE: PERL Program
SOFTWARE: PERL Program
SEQ ID NO 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 29, Application US/09919039 Publication No. US20030108871A1 GENERAL INFORMATION:
                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Kaser, Matthew R.
TITLE OF INVENTION: GENES EXPRESSED IN TREATED HUMAN C3A LIVER CELL CULTURES
FILE REFERENCE: PA-0035 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: CuraSeqList version 0.1
                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 98.5%; Score 660; DB 5; ]
Local Similarity 100.0%; Pred. No. 3.2e-61;
les 123; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          284 AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
                               AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYFTVNCQP 122
                                                                                                                            GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
                                                                                                GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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                                                                                                                                                                                                98.5%; So ilarity 100.0%; I Conservative 0;
                                                                                                                                                                                                                                                                                                                               No.
                                                                                                                                                                                                                          Score 660; DB 3; ; Pred. No. 3.5e-61;
                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                 Length 427;
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                                                                                                     312
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; LENGTH: 427
; TYPE: PRT
; ORGANISM: Homo 9
US-10-637-313-24
RESULT 11
US-10-741-600-1179
; Sequence 1179, Application US/10741600
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CURRENT APPLICATION NUMBER: US/10/637,313
CURRENT FILING DATE: 2003-08-08
PRIOR APPLICATION NUMBER: 10/162335
PRIOR APPLICATION NUMBER: 60/295607
PRIOR APPLICATION NUMBER: 60/295661
PRIOR APPLICATION NUMBER: 60/295661
PRIOR FILING DATE: 2001-06-04
PRIOR PILLING DATE: 2001-06-04
PRIOR PILLING DATE: 2001-06-06
PRIOR PILLING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: 60/29828
PRIOR APPLICATION NUMBER: 60/29828
PRIOR APPLICATION NUMBER: 60/298949
PRIOR PILLING DATE: 2001-06-21
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US-10-637-313-24
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APPLICANT: Uriel M
APPLICANT: Li Li
APPLICANT: David A
APPLICANT: Sasha C
APPLICANT: Mei Zho
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Best Local Similarity
Matches 123; Conserva
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Publication No. US20040229779A1
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/300883
PRIOR FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: 60/301550
PRIOR FILING DATE: 2001-06-28
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TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHOD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                         63 AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYFTVNCQP 122
                                                                                                                                                                    LGM 125
                                                                                                                                                                                                                                                                                                                                 GKDFVQPFTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV 62
                                                                                                                                                                                                                 AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
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Nikolai Khramtsov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Stacie Casman
Ferenc Boldog
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RESULT 13
US-10-893-315-101
; Sequence 101, Application US/10893315
; Publication No. US20050147987A1
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RESULT 12

US-10-741-600-1181

US-10-741-600-1181

Sequence 181, Application US/10741600

Publication No. US20050026169A1

GENERAL INFORMATION:
APPLICANT: CARGILL, Michele et al.
TITLE OF INVENTION: MYOCARDIAL INFARCTION, METHODS OF DETECT FILE REFERENCE: CL001499

CURRENT APPLICATION NUMBER: US/10/741,600

CURRENT FILING DATE: 2003-12-22

NUMBER OF SEQ ID NOS: 73997

SOFTWARE: FASCUSE OF Windows Version 4.0

SEQ ID NO 1181

LENGTH: 427

TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Publication No. US20050026169A1
; GENERAL INFORMATION:
APPLICANT: CARGILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WIT
TITLE OF INVENTION: MYOCARDIAL INFARCTION, METHODS OF D
FILE REFERENCE: CL001499
; CURRENT APPLICATION NUMBER: US/10/741,600
; CURRENT FILING DATE: 2003-12-22
; NUMBER OF SEQ ID NOS: 73997
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1179
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-741-600-1179
                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Homo sapiens 
US-10-741-600-1181
                                                                                                                                                                                                                                                                                                              Query Match
Best Local
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Local Similarity 100.0%; Pred. No. 3.5e-61;
hes 123; Conservative 0; Mismatches 0;
373
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                                                                                                                                                                                            253 GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
                                                                                                                               63 AGKKYPIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP 122
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375
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100.0%; Pred. No. 3.5e-61;
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GEI
TITLE OF INVENTION: WITH TYPE II DIABETES ANI
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL000786
CURRENT APPLICATION NUMBER: US/10/893,315
CURRENT FILING DATE: 2004-07-19
PRIOR APPLICATION NUMBER: 60/231,397
PRIOR APPLICATION NUMBER: 60/231,397
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 2172
SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 105
LENGTH: 579
TYPE: PRT
ORGANISM: Human
US-10-893-315-105
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TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GE!
TITLE OF INVENTION: WITH TYPE II DIABETES AN!
TITLE OF INVENTION: USES THEREOF
FILE REFERENCE: CL000786
CURRENT APPLICATION NUMBER: US/10/893,315
CCURRENT FILING DATE: 2004-07-19
PRIOR APPLICATION NUMBER: 60/231,397
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 2172
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 101
LENGTH: 579
TYPE: PRT
ORGANISM: Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 105, Application US/10893315 Publication No. US20050147987A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 123; Conserv
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                                                                                                                                                                                  GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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                                     LGM 125
                                                                          AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP
                                                                                                   AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP 122
                                                                                                                                                       GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
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TGM 310
                                                                                                                                                                                                                                  98.5%; Score 660; DB 5; I ilarity 100.0%; Pred. No. 5.2e-61; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                          Length 579;
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APPLICANT: ZOTHUBEN, Bryan D.
APPLICANT: ZOTHUBEN, Bryan D.
TITLE OF INVENTION: Therapeutic Polypeptides, Nucleic Acids Encoding Same, and Method
FILE REFERENCE: 21402-377 B
FILE REFERENCE: 21402-377 B
CURRENT APPLICATION NUMBER: US/10/162,335
CURRENT FILING DATE: 2002-10-01
PRIOR APPLICATION NUMBER: 60/295,607
PRIOR FILING DATE: 2001-06-04
PRIOR FILING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: 60/295,661
PRIOR APPLICATION NUMBER: 60/296,404
PRIOR APPLICATION NUMBER: 60/296,404
PRIOR APPLICATION NUMBER: 60/296,418
PRIOR APPLICATION NUMBER: 60/296,418
PRIOR APPLICATION NUMBER: 60/297,414
PRIOR APPLICATION NUMBER: 60/297,414
PRIOR PILING DATE: 2001-06-06
PRIOR APPLICATION NUMBER: 60/297,567
PRIOR APPLICATION NUMBER: 60/297,567
PRIOR APPLICATION NUMBER: 60/297,567
PRIOR APPLICATION NUMBER: 60/297,567
PRIOR APPLICATION NUMBER: 60/290,567
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US-10-162-335-72
US-10-162-335-72
; Sequence 72, Application US/10162335
; Publication No. US20040009480A1
                                                                                                                                                                                                                                                     US-10-162-335-72
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NUMBER OF SEQ ID NOS: 201
SEQ ID NO 72
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                                                                                                                                                      Query Match 98.5%; Score 660; DB 4; Best Local Similarity 100.0%; Pred. No. 5.6e-61; Matches 123; Conservative 0; Mismatches 0;
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TYPE: PRT
ORGANISM: Homo
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                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/300,883 PRIOR FILING DATE: 2001-06-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/298,556
PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/299,949
PRIOR FILING DATE: 2001-06-21
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                                                                   GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
                                                                                              GKDFVQBPTKICVGCBRDIPTNSBELEETLTHTITKLNAENNATFYFKIDNVKXARVQVV
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Pena, Carol E. A.
Rastelli, Luca
Shimkets, Richard A.
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Guo, Xiaojia (Sasha)
Hjalt, Tord
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Gerlach, Valerie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Spytek, Kimberly A. Vernet, Corine A. M. Voss, Edward Z.
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Malyankar, Uriel M.
Millet, Isabelle
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Casman, Stacie J.
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                                                                                                                                                                                                                                                                                                                                                                                    data removed -
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                                                                                                                                                                                                    Length 615;
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Qy 123 LGM 125 Db 344 LGM 346 Search completed: February 8, 2006, 03:21:54 Job time: 179 secs	284 AGKKYFIDEVARRTTCSKESNEELTESCETKKLGOSLDCNARVYVBWEKKTVDTUNCOD 343

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Minimum
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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1: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

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670
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Copyright (c) 1993 - 2006 Biocceleration Ltd.
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SULT 2 -10-967- Sequence Publicat GENERAL APPLICA TITLE O			Query M Best Lo Matches	Sequence: Sequence: Sequence: Sequence: GENERAL III APPLICAN INIMA APPLICAN INIMA APPLICAN AP		22222222222222222222222222222222222222
SULT 2 -10-967-093-8 Sequence 8, Application US/10: Sequence 10, Application US/10: SENERAL INFORMATION: APPLICANT: Schiemann, Willian TITLE OF INVENTION: Cystation FILE REFERENCE: 2879-104	71 FV. 114 VE	11 TK :: 54 SR	atch cal	197, 197, INFORMATION INTO THE STATE OF TH		61.5 61.5 61.5 61.5 61.5 61.5 61.5 61.5
plication . . US200502 ATION: hiemann, W NTION: Cys	FVARETTCSKESNEELTESCE- : : : VEIGRTTCKKNQHLRL-DDCDF	TKICVGCPRDIPTNS ::: : SRVKPGFPKTIKTND	Similarity 7; Conservat	Application US/1068974 Application US/1068974 O. US20050250180A1 MATION: acobs, Kenneth MCCOy, John M LaVallie, Edward R Racie, Lisa A Rvans, Cheryl Merberg, David Treacy, Maurice Spaulding, Vikki ENTION: SECRETED PROTEI CB: 00766.000091.10 ICATION NUMBER: US/10/6 MG DATE: 2003-10-22 ATION NUMBER: 09/746,78 DATE: 2000-12-21 Q ID NOS: 231 tentin version 3.2 como sapiens		88889999999999999999999999999999999999
on US/109 50267021A William Cystatin	CESNEE : CNQHLR	DIPTNS KTND	24. 32. ive	sion U so2501 bhu ma bhu ma bha m bha m bha m bha m bha m bha m bha m bha m sion m sio		1038 6284 6285 1178 1178 2053 2066 227 1306 334 391 407 603 603 651 193 193 248
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. P. C as an Antagonist of TGF-	FVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKI-YPTVNC 	TKI CVGCPRDI PTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKYFID :::	Score 165.5; DB 6; L Pred. No. 1.6e-10; 22; Mismatches 50; I	0689742 1 R PROTEINS AND POLYNUCLEOTIDE 5/10/689,742 -22 746,783	ALIGNMENTS	US-11-107-028-1 US-11-075-351-12 US-10-793-626-2844 US-11-044-899-29 US-10-877-346-38 US-10-877-346-91 US-10-877-346-91 US-10-877-346-91 US-10-877-346-91 US-10-485-517-210 US-10-485-517-241 US-10-485-517-241 US-11-016-564-1 US-11-016-564-7 US-11-1016-564-7 US-11-183-136-2 US-11-183-136-2 US-11-183-136-2 US-11-183-136-2 US-11-183-136-2 US-11-183-136-2 US-11-183-136-2 US-11-183-136-3 US-10-928-360A-558 US-10-928-360A-558 US-10-955-561-959 US-11-057-058-59
F-B and Methods Related Thereto	WEKKI-YPTVNC 120 :	NVKKARVQVVAGKKYFID 70 : : RITRALVQIVKGLKYMLE 113	Length 167; Indels 5; Gaps 3;	IDES ENCODING THEM		Sequence 1, Appli Sequence 12, Appl Sequence 29, Appl Sequence 29, Appl Sequence 11, Appl Sequence 11, Appl Sequence 210, Appl Sequence 210, App Sequence 211, Appl Sequence 241, Appl Sequence 241, Appl Sequence 1, Appli Sequence 1, Appli Sequence 7, Appli Sequence 7, Appli Sequence 300, Appl Sequence 300, Appl Sequence 558, Appl Sequence 1666, Ap Sequence 559, Appl

FILE REFERENCE: 2879-104

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Sequence 6, Application US/10967093
Publication No. US20050267021A1
GENERAL INFORMATION:
APPLICANT: Schiemann, William P.
TITLE OF INVENTION: Cystatin C as an Antagonist of FILE REFERENCE: 2879-104
CURRENT APPLICATION NUMBER: US/10/967,093
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: 60/511,794
PRIOR APPLICATION NUMBER: 60/511,794
PRIOR APPLICATION NUMBER: 60/511,794
PRIOR FILING DATE: 2003-10-15
NUMBER OF SEQ ID NOS: 10
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Matches
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SEQ ID NO 8
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SOFTWARE: PatentIn version 3.1 SEQ ID NO 6
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Best Local Similarity
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-10-967-093-8
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TITLE OF INVENTION: Cystatin C as an Antagonist of TGF-B and Methods Related Thereto FILE REFERENCE: 2879-104
CURRENT APPLICATION NUMBER: US/10/967,093
CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: 60/511,794
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NUMBER OF SEQ ID NOS:
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CURRENT FILING DATE: 2004-10-15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    106 QANLDSCPFHNQPHLKREKL----CSFQVYVVPWMN----TINLVKFSCQ 147
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                                                                                                                                                                               of TGF-B and Methods Related Thereto
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Sequence 425, Application US/09978360A

Publication No. US20060009633A9

GENERAL INFORMATION:

APPLICANT: Edwards, Jean-Baptiste Dumas Milne

APPLICANT: Duclert, Aymeric

APPLICANT: Dobert, Severin

APPLICANT: Clusel, Catherine

FITLE OF INVENTION: Complementary DNA's Encoding Proteins with Signal Peptides

FILE REFERENCE: 56.US4.CIP

CURRENT APPLICATION NUMBER: US/09/978,360A

CURRENT APPLICATION NUMBER: US 60/066,677

PRIOR FILING DATE: 1997-11-13

PRIOR APPLICATION NUMBER: US 60/069,957

PRIOR APPLICATION NUMBER: US 60/069,957

PRIOR APPLICATION NUMBER: US 60/069,957

PRIOR APPLICATION NUMBER: US 60/069,957
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US-10-967-093-4
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Publication No. US20050267021A1
GENERAL INFORMATION:
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Best Local Similarity
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CURRENT FILING DATE: 2004-10-15
PRIOR APPLICATION NUMBER: 60/511,794
PRIOR FILING DATE: 2003-10-15
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Schiemann, William P. TITLE OF INVENTION: Cystatin C as an Antagonist of TGF-B and Methods Related Thereto FILE REFERENCE: 2879-104
PRIOR APPLICATION NUMBER: US 60/074,121
PRIOR FILING DATE: 1998-02-09
PRIOR APPLICATION NUMBER: US 60/081,563
PRIOR FILING DATE: 1998-04-13
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Local Similarity 30.0%;
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US-09-978-360A-425
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Best Local Similarity
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SEQ ID NO 425
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Publication No. US20050024400000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: McGuire, James
APPLICANT: Simonsen, Anja Hviid
APPLICANT: Blennow, Kaj
APPLICANT: Podust, Vladinir
APPLICANT: Ciphergen Biosystems, Inc.
TITLE OF INVENTION: Biomarkers for Alzheimer's Disease
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                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 016866-011550US
CURRENT APPLICATION NUMBER: US/10/982,545
CURRENT FILING DATE: 2004-11-06
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PRIOR
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APPLICANT: McGuire, James
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                                                                                                        NUMBER OF SEQ ID NOS:
                                                                                                                             PRIOR APPLICATION NUMBER: US PRIOR FILING DATE: 2004-07-08
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo Sapiens
                                                ENGTH: 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH: 382
                                                                                                                                                                                                       APPLICATION NUMBER: US 60/546,423
FILING DATE: 2004-02-19
APPLICATION NUMBER: US 60/547,250
FILING DATE: 2004-02-23
APPLICATION NUMBER: US 60/558,896
FILING DATE: 2004-04-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 1998-12-17
APPLICATION NUMBER: US 09/247,155
FILING DATE: 1999-02-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 09/215,435
FILING DATE: 1998-12-17
APPLICATION NUMBER: PCT/IB98/02122
                                                                                                                                                                 APPLICATION NUMBER: US 60/572,617 FILING DATE: 2004-05-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US 09/191,997 FILING DATE: 1998-11-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 60/099,273 FILING DATE: -09-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 60/096,116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           210 LIKESPCTKSQ----ASSCSL----QSSDSVPVGLCKGSLTRTHWEK--FVSVTC 254
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                                                                                        Patentin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VARETTCSKESNEELTESCETKKLGQSLD-----CNAEVYVVPWEKKIYPTVNC 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CVGCPRDIPTNSP--ELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKYFIDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19.3%; Score 129.5; DB 5 ilarity 30.4%; Pred. No. 2.6e-06; Conservative 18; Mismatches 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1998-08-10
                                                                                                                           2004-07-08
                                                                                                                                                60/586,503
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RESULT 9
US-10-821-234-1237
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Best Local Similarity
Matches 30; Conserv
                                                                                                       Sequence 1237, Application US/10821234 Publication No. US20050255114A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/10967093
Publication No. US20050267021A1
GENERAL INFORMATION:
APPLICANT: Schiemann, William P.
TITLE OF INVENTION: Cystatin C as an Antagonist of TGF-B and Methods Related Thereto
FILE REFERENCE: 2879-104
CURRENT APPLICATION NUMBER: US/10/967,093
CURRENT FILING DATE: 2004-10-15
PRICE REFERENCE: 2004-10-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn version 3.1 SEQ ID NO 2
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                                  APPLICANT:
                                                                     APPLICANT: Labat, Ivan
APPLICANT: Stache-Crain, Birgit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/511,794 PRIOR FILING DATE: 2003-10-15 NUMBER OF SEQ ID NOS: 10
             APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis
FILE REFERENCE: 821A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: PEPTIDE
LOCATION: (35)..(146)
OTHER INFORMATION: biomarker peptide M12583.4, truncated Cystatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
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LOCATION: (1)..(26)
OTHER INFORMATION:
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LOCATION: (27)..(146)
OTHER INFORMATION: biomarker peptide M13391, full-length Cystatin
OTHER INFORMATION: mature peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Cystatin C precursor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           89
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   146
                                                                                                                                                                                                                                                                                                                                            8 QPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKY
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                                                                                                                                                                                                                                                                       FIDFVARETTCSKESNEELTESC---ETKKLGQSLDCNAEVYVVPWE 111
                                                                                                                                                                                                                                    FLDVELGRTTCTK--TOPNLDNCPFHDOPHLKRKAFCSFQIYAVPWQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FLDVELGRTTCTK--TQPNLDNCPFHDQPHLKRKAFCSFQIYAVPWQ 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FIDFVARETTCSKESNEELTESC---ETKKLGQSLDCNAEVYVVPWE 111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVVAGKKY
                                                     Andarmani,
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28.0%; Pref
-tve 21;
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Pred. No. 1.3e
21; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                    Score 127.5; DB 6;
pred. No. 1.3e-06;
1; Mismatches 49;
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                 and Treatment
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                                                                                                                                                                                                                                                                                                                                                                                                                       Length 146;
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                 of Preeclampsia
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Sequence 8, Application US/10888962
Publication No. US20050266531A1
GENERAL INFORMATION:
APPLICANT: Saint Louis University
APPLICANT: Ray, Ranjit
APPLICANT: Yie-Hwa, Chang
APPLICANT: Ray, Ratna
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                                                                                                                                                 US-10-888-962-8
                                                                                                                                                                  RESULT 11
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Best Local S
Matches 25
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Best Local S
Matches 24
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CURRENT FILING DATE: 2004-04-07
PRIOR APPLICATION NUMBER: US 60/462,047
PRIOR FILING DATE: 2003-04-07
NUMBER OF SEQ ID NOS: 1704
SOFTWARE: pt SEQ_genes Version 1.0
SEQ ID NO 1237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 7, Application US/10888962 Publication No. US20050266531A1 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Saint Louis University
APPLICANT: Ray, Ranjit
APPLICANT: Yie-Hwa, Chang
APPLICANT: Ray, Ratna
APPLICANT: Basu, Arnab
TITLE OF INVENTION: Compositions and Methods for Inhibiting Liver Stellate Cell
TITLE OF INVENTION: Growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: SLU 03-013 PCT
CURRENT APPLICATION NUMBER: US/10/888,962
CURRENT FILING DATE: 2004-07-09
PRIOR APPLICATION NUMBER: 60/487,126
PRIOR FILING DATE: 2003-07-12
NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 345
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                                                                                                                                                                                                                              192 PLPVSTLVEFVIAATDCTA---KEVTDPAKCNLLAE 224
                                                                                                                                                                                                                                                                                                                134
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Similarity 28.9%; Pred. No. 4.6e-05;
24; Conservative 19; Mismatches 35
                                                                                                                                                                                                                                                                                                                                                                                               Similarity 26.(
25; Conservative
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                                                                                                                                                                                                                                                                       VAGKKYFIDFVARETTCSKESNEELTESCETKKLGQ 97
                                                                                                                                                                                                                                                                                                                STPDSAEDVRKLCPRCPLLTPFNDTNVVHTVNTALAAFNTQNNGT-YFKLVEISRAQ-NV 191
                                                                                                                                                                                                                                                                                                                                                                                                                14.1%; Score 94.5; DB 6; Length 345; 26.0%; Pred. No. 0.01;
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FILE REFERENCE: SLU 03-013 PCT; CURRENT APPLICATION NUMBER: US/10/888,962; CURRENT FILING DATE: 2004-07-09; PRIOR APPLICATION NUMBER: 60/487,126; PRIOR FILING DATE: 2003-07-12; NUMBER OF SEQ ID NOS: 15; SOFTWARE: PALENTIN VERSION 3.2; SEQ ID NO 8
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SEQ ID NO 6
LENGTH: 367
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Best Local Similarity
Matches 26; Conserv
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Publication No. US20050266531A1
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Best Local
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APPLICANT: Ray, Ranjit
APPLICANT: Yie-Hwa, Chang
APPLICANT: Ray, Ratna
APPLICANT: Basu, Arnab
TITLE OF INVENTION: Crowth
TITLE OF INVENTION: Growth
FILE REFERENCE: SLU 03-013 PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 60/487,126
PRIOR FILING DATE: 2003-07-12
NUMBER OF SEQ ID NOS: 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/10/888,962
CURRENT FILING DATE: 2004-07-09
                                                                                                                                                                                                                                                                                                                                                              LENGTH: 36
TYPE: PRT
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ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                           Match 13.2%; Score 88.5; DB 6; Local Similarity 26.1%; Pred. No. 0.048;
                                                                                                                                                       134 SSPDSAEDVRKYCQDCPLLAPLNDTRVVHAAKAALAAFNAQNNGS-NFQLEEISRAQLVP 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    192 PFPVSTLVEFVIAATDC---TGQEVTDPAKCNLLAE 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  134 STPDSAEDVRKFCPRCPILIRENDTNVVHTVKTALAAFNAQNNGT-YFKLVEISRAQ-NV 191
247
                                                                           193 LPPSTY-VEFTVSGTDCVAKEATEAAKCNLLAEKQYGFCKATLSEKLG---
                                     109 PWEKKIYPTVNCQP 122
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                                                                                                                 62 VAGKKYFIDFVARETTC-SKESNEE-----LTES----CE---TKKLGQSLDCNAEVYVV 108
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CTVFQTQP-VTSQP
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259
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RESULT 13 US-10-467-962B-12 ; Sequence 12, Apr

Application US/10467962B

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APPLICANT: Plach, Gunnar
APPLICANT: Blau, Astrid
APPLICANT: Daeschner, Klaus
APPLICANT: Daeschner, Klaus
APPLICANT: Klein, Mathieu
ITITLE OF INVENTION: Identification of Herbicidally Active Substances
FILE REFERENCE: 2000 857
CURRENT APPLICATION NUMBER: US/10/467,962B
CURRENT FILING DATE: 2003-08-14
PRIOR APPLICATION NUMBER: PCT/EP02/01466
PRIOR APPLICATION NUMBER: PCT/EP02/01466
PRIOR FILING DATE: 2002-02-13
NUMBER OF SEQ ID NOS: 109
SOFTWARE: Patentin Vers. 2.0
SEQ ID NO 12
LENGTH: 758
                                                                                              ; ORGANISM: Homo sapiens
US-11-137-465-37
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US-11-137-465-37
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                                                                                                                                                                                                                                            CURRENT FILING DATE: 2005-05-25
PRIOR APPLICATION NUMBER: US/10/239,663
PRIOR APPLICATION NUMBER: PCT/US01/09226
PRIOR APPLICATION NUMBER: PCT/US01/09226
PRIOR APPLICATION NUMBER: 60/192,158
PRIOR APPLICATION NUMBER: 60/192,158
PRIOR PILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: 60/192,668
PRIOR FILING DATE: 2000-03-27
PRIOR PILING DATE: 2000-03-27
PRIOR APPLICATION NUMBER: 60/200,166
PRIOR APPLICATION NUMBER: 60/200,166
PRIOR FILING DATE: 2000-03-27
PRIOR FILING DATE: 2000-04-27
                                                                                                                                      SEQ ID NO 37
LENGTH: 138
TYPE: PRT
  Query Match 11.0%;
Best Local Similarity 24.5%;
Matches 23; Conservative 2
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Agarwal, Pankaj
APPLICANT: Murdoch, Paul
APPLICANT: Rizvi, Safia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 66
SOFTWARE: FastSEQ for Wil
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CURRENT APPLICATION NUMBER: US/11/137,465
CURRENT FILING DATE: 2005-05-25
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TITLE OF INVENTION: NOVEL COMPOUNDS
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                                                                                                                                                              138
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Rizvi, Safia, K.
Smith, Randall, F.
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20;
  Score 74; DB
Pred. No. 0.47
20; Mismatches
                       DB 7; Length 138; 0.47;
47; Indels
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; SEQ ID NO 5
; LENGTH: 359
; TYPE: PRT
; ORGANISM: BOB taurus
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                                                                                                                                                                                                                                                                                                                                               FILE REFERENCE: SLU 03-013 PCT CURRENT APPLICATION NUMBER: US/10/888,962 CURRENT FILING DATE: 2004-07-09 PRIOR APPLICATION NUMBER: 60/487,126 PRIOR FILING DATE: 2003-07-12 NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Basu, Arnab
TITLE OF INVENTION: Compositions and Methods for Inhibiting Liver Stellate Cell
TITLE OF INVENTION: Growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Saint Louis University
APPLICANT: Ray, Ranjit
APPLICANT: Yie-Hwa, Chang
                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn version 3.2
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                                                                                                                                                                        Local Similarity
  190 ---
                                                                        134 SSPDSAEDVRKLCPDCPLLAPLNDSRVVHAVEVALATFNAESNGS-YLQLVEISRAQ---
                                  62 VAGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCN 102
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                                                                                                                                                      23;
                                                                                                              2 SGKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQV 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/10888962
No. US20050266531A1
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                                                                                                                                                      Conservative
- FVPLPVSVSVEFAVAATD-CIAKEVVDPTKCN
                                                                                                                                                  10.7%; Score 72; DB 22.8%; Pred. No. 2.5; Ative 16; Mismatches
                                                                                                                                                                                        DB 6; Length 359;
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Search completed: February 8, 2006, 03:22:16
Job time: 17 secs

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     GenCore version (c) 1993 - 2006
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T-kininogen, LMW I
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12.6	13.2	13.2	13.2	13.2	13.7	13.7	13.7	14.0	14.1	14.2	14.3	14.5	14.6	15.7
592	367	134	134	125	375	135	100	139	345	142	143	133	135	162
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T50516	WOHU	S54828	JC4882	T00752	A32827	S27239	JH0269	A45361	S21094	A47142	T33301	JC4536	JC4007	A43428
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Ca2+ dependent lip	alpha-2-HS-glycopr	cysteine proteinas	cystatin - maize	cysteine proteinas	fetuin precursor -	cysteine proteinas	cystatin - avocado	cystatin-related e	alpha-2-HS-glycopr	cystatin D precurs	hypothetical prote	cystatin precursor	cystatin II - maiz	onchocystatin - ne

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A;Molecule type: protein A;Residues: 381-389 <mae> A;Cross-references: UNIPARC:UPI000002CF4A A;Cross-references: UNIPARC:UPI000002CF4A R;Sasaguri, M.; Ikeda, M.; Ideishi, M.; Arakawa, K. R;Sasaguri, M.; Ikeda, M.; Ideishi, M.; Arakawa, K. Biochem. Biophys. Res. Commun. 150, 511-516, 1988 A;Title: Identification of [hydroxyproline(3)]-lysyl-bradykinin A;Reference number: A34030; MUID:88106632; PMID:3337729</mae>	A; Molecule type: protein A; Residues: 380-389 < MIN> A; Cross-references: UNIPARC: UPI000002COF8 A; Cross-references: UNIPARC: UPI000002COF8 R; Maeda, H.; Matsumura, Y.; Kato, H. J. Biol. Chem. 263, 16051-16054, 1988 A; Title: Purification and identification of [hydroxyproly1(3)]bradykinin in ascitic A; Reference number: A31905; MUID:89034061; PMID:3182782 A; Accession: A31905	A;ACCESSION: A2/900 A;Molecule type: protein A;Residues: 390-427 <lot> A;Residues: 390-427 <lot> A;Cross-references: UNIPARC:UPI00001742A0 A;Cross-references: UNIPARC:UPI00001742A0 B;Cross-references: UNIPARC:UPI00001742A0 CA:, Proud, D:; Walz, D:; 1988 Biochem. Biophys. Res. Commun. 152, 519-526, 1988 A;Title: A new kinin moiety in human plasma kininogens A;Reference number: A27699; MUID:88209021; PMID:336523 A;Accession: A27699</lot></lot>	A; Molecule type: mENN A; Residues: 1-427 <tak> A; Residues: 1-427 <tak> A; Residues: 1-427 <tak> A; Residues: 1-427 <tak> A; Cross-references: UNIFARC: UPI00000D92C; GB:M11437; NID:g186751; PIDN:AAB59551.1; PI A; Cross-references: UNIFARC: UPI00000D92C; GB:M11437; NID:g186751; PIDN:AAB59551.1; PI R; Lottspeich, F.; Kellermann, J.; Henschen, A.; Rauth, G.; Mueller-Esterl, W. in Kinins IV, part A, Greenbaum, L.M., and Margolius, H.S., eds., pp.91-95, Plenum, Ne A; Title: Amino acid sequence of the light chain of human low molecular mass kininogen. A; Reference number: A27900</tak></tak></tak></tak>	A;Molecule type: mRNA A;Residues: 1-427 cOHK> A;Residues: 1-427 cOHK> A;Cross-references: UNIPROT:P01042; UNIPARC:UPI000000D92C; GB:K02566; NID:g177889; PIDN A;Cross-references: UNIPROT:P01042; UNIPARC:UPI000000D92C; GB:K02566; NID:g177889; PIDN R;Takagaki, Y.; Kitamura, N.; Nakanishi, S. J. Biol. Chem. 260, 8601-8609, 1985 J. Biol. Chem. 260, 8601-8609, 1985 A;Title: Cloning and sequence analysis of cDNAs for human high molecular weight and low A;Reference number: A92544; MUID:85234582; PMID:2989293	RESULT 1 KGHUL1 KININOGEN, LAW precursor [validated] - human kininogen, Law precursor [validated] - human N;Alternate names: alpha-2-thiol proteinase inhibitor; preprokininogen N;Contains: bradykinin (kallidin); kininogen I; kininogen II; prokininogen C;Species: Homo sapiens (man) C;Date: 06-Jul-1982 #sequence revision 27-Nov-1985 #text change 09-Jul-2004 C;Accession: A01280, B25276; A27900; A27699; A31905; A34030 R;Ohkubo, I.; Kurachi, K.; Takasawa, T.; Shiokawa, H.; Sasaki, M. Biochemistry 23, 5691-5697, 1984 A;Title: Isolation of a human cDNA for alpha-2-thiol proteinase inhibitor a A;Reference number: A90490; MUID:85122621; PMID:6441591 A;Accession: A01280		39 91.5 13.7 375 2 40 88.5 13.2 13.5 2 41 88.5 13.2 13.4 2 42 88.5 13.2 13.4 2 43 88.5 13.2 13.4 2 44 85 12.7 199 2 45 84.5 12.6 592 2
)0002CF4A 11, M.; Arakawa, K. 50, 511-516, 1988 xxyproline(3)]-lysyl-bradyk :88106632; PMID:3337729)0002C0F8 H. 1988 Eleation of [hydroxyprolyl(0001742A0 Proud, D.; Walz, D.; Scicli, 52, 519-526, 1988 Iman plasma kininogens. 188209021; PMID:3365237	0000D92C; GB:M11437; NID:g; Henschen, A.; Rauth, G.; L.M., and Margolius, H.S., L.M., the light chain of human lo	12; UNIPARC:UPI000000D92C; canishi, S. 985 1986 of cDNAs for human h	proteinase inhibitor; prep proteinase inhibitor; prep), kininogen I; kininogen I /ision 27-Nov-1985 #text ch 900; A27699; A31905; A34030 awa, T.; Shiokawa, H.; Sasa, wa for alpha-2-thiol protei	ALIGNMENTS	A32827 T00752 JC4882 S54828 WOHU S65071 T50516
inin released from human plas	3)]bradykinin in ascitic fluı	i, A.G.	186751; PIDN:AAB59551.1; PID Mueller-Esterl, W. eds., pp.91-95, Plenum, New w molecular mass kininogen.	GB:K02566; NID:g177889; PIDN	rokininogen I; prokininogen ange 09-Jul-2004 ki, M.		fetuin precursor - cysteine proteinas cystatin - maize cysteine proteinas alpha-2-HS-glycopr cystatin - field m ca2+ dependent lip

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A,Gene: GDB: KNG
A,Gene: GDB: KNG
A,Gross-references: GDB:125256; OMIM:228960
A;Map position: 3q27-3q27
A;Map position: 3q27-3q27
A;Introns: 65/3; 102/3; 131/1; 188/3; 224/3; 253/1; 310/3; 346/3; 375/3; 401/3
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F;389-390/Cleavage site: Met-Lys (Kallikrein) #status experimental
F;389/390/Cleavage site: Ang-Sear (Kallikrein) #status experimental
F;401/Binding site: carbohydrate (Thr) (covalent) #status experimental
F;401/Binding site: carbohydrate (Thr) (covalent) #status experimental
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A;Title: Structural organization of the human kininogen gene and a A;Reference number: A92545; MUID:85234583; PMID:2989294
A;Contents: annotation; gene organization
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A;Molecule type: protein
A;Residues: 380-389 <SAS
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C;Date: 28-May-1986 #sequence revision 28-May-1986 #text_change 09-Jul-2004
C;Date: 28-May-1986 #sequence revision 28-May-1986 #text_change 09-Jul-2004
C;Accession: A01279; AS5276; S32422; A91153; A24871; A77899; A27699; A31905
R;Ohkubo, I.; Kurachi, K.; Takasawa, T.; Shiokawa, H.; Sasaki, M.
Biochemistry 23, S691-5697, 1984
Biochemistry 23, S691-5697, 1984
A;Title: Isolation of a human cDNA for alpha-2-thiol proteinase inhibitor a:
A;Reference number: A90490; MUID:85122621; PMID:6441591
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                                                                                                                                                                                                                                          KGHUH1
kininogen, HMW precursor [validated] - human
kininogen, HMW precursor [validated] - human
kininogen inhibitor;
N;Contains: bradykinin (kallidin I); HMW kininogen I; H
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100.0%; Pred. No. 7.8e-55;
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HMW kininogen II;
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low molecular
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A;ACCCEPTON.
A;Molecule type: protein
A;Molecule type: protein
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A;Residues: 380-389 <MIN>
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R;Maeda, H.; Matcumura, Y.; Kato, H.
R;Maeda, H.; Matcumura, Y.; Kato, H.
J. Biol. Chem. 263, 16051-16054, 1988
A;Title: Purification and identification of [hydroxyprolyl(3)]bradykinin
A;Title: Purification and identification of [hydroxyprolyl(3)]bradykinin
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A; Molecule type: protein
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A; Cross-references: UNIPARC: UPI000017429A
A; Cross-references: UNIPARC: Proud, D.; Walz, D.; Sc
R; Mindrou, T.; Carretero, O.A.; Proud, D.; Walz, D.; Sc
Biochem. Biophys. Res. Commun. 152, 519-526, 1988
Biochem. Biophys. Res. Commun. 152, 519-526, 1988
A; Title: A new kinin molety in human plasma kininogens.
A: Reference number: A27699; MUID:88209021; PMID:3365237
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A; Reference number: S32422; MUID:9323854; PMID:8467916
A; Accession: S32422
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A; Residues: 'ANSM', 253-377 < AUE>
A; Cross-references: UNIPARC:UPI0000174297
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A; Mote: differences are due to known cloning artifacts
A; Mote: differences are due to known cloning artifacts
A; Cross-references are due to known cloning artifacts
A; Mote: differences are due to known cloning artifacts
A; Mote: difference number: MIPARC:UPI000174297
A; Mote: difference number: 152, 307-314, 1985
A; Mille: The amino acid sequence of the light chain of human high-molecular-mass
A; Meference number: A91153; MUID:86030270; PMID:4054110
A; Accession: A91153
A; Molecule type: protein
A; Residues: 379-644 CLOTY
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A;Cross-references: UNIPROT:P010429; UNIPROT
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A;Note: the bradykinin sequence preceding the light chain sequence R;Kellermann, J.; Lottspeich, F.; Henschen, A.; Mueller-Esterl, W. Eur. J. Biochem. 154, 471-478, 1986
Eur. J. Biochem. 154, 471-478, 1986
A;Title: Completion of the primary structure of human high-molecula A;Reference number: A24871; MUID:86108361; PMID:3484703
A;Accession: A24871
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J. Biol. Chem. 260, 8601-8609, 1985
A;Title: Cloning and sequence analysis of cDNAs for human high molecular weight and A;Reference number: A92544; MUID:85234582; PMID:2989293
A;Accession: A25276
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A;Cross-references: UNIPARC:UPI000002CF4A
A;Cross-references: UNIPARC:UPI000002CF4A
A;Rakawa, K.
Biochem, Biophys. Res. Commun. 150, 511-516, 1988
Biochem, Biophys. Res. Commun. 150, 511-516, 1988
A;Title: Identification of [hydroxyproline(3)]-lysyl-bradykinin released from A;Reference number: A34030; MUID:88106632; PMID:3337729
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A;Cross-references: UNIPARC:UPI0000174299
R;Lenarcic, B.; Gabrijelcic, D.; Rozman, B.; Drobnic-Kosorok, Biol. Chem. Hoppe-Seyler 369, 257-261, 1988 A;Tille: Human cathepsin B and cysteeine proteinase inhibitors A;Reference number: S02482; MUID:89076517; PMID:3264507
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PMID:3484703
                                                                                                                                                                                        Drobnic-Kosorok, M.; Turk,
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A;Cross-references: GDB:125256; OMIM:228960
A;Map position: 3427-3427
A;Introns: 65/3; 102/3; 131/1; 188/3; 224/3; 253/1; 310/3; 346/3; 375/3
A;Introns: 65/3; 102/3; 131/1; 188/3; 224/3; 253/1; 310/3; 346/3; 375/3
C;Superfamily: kininogen; cystatin homology
C;Keywords: alternative splicing; blood coagulation; cysteine proteinase inhibitor; dupl
F;1-18/Domain: signal sequence #status experimental <SIG>
F;19-644/Product: HMW kininogen I (prokininogen) #status experimental <MATI>
F;19-379,390-644/Product: HMW kininogen II #status experimental <MATI>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Comment: The HMW kininogen precursor and the LMW form are produced from the same gene C;Comment: Kininogen is a cysteine proteinase inhibitor, takes part in initiation of the C;Comment: The glycine/histidine/lysine-rich region of HMW kininogen light chain is impor C;Comment: Bradykinin, released from kininogen by kallikrein, is a potent vasodilator, in a potent vasodilator, in a potent vasodilator, in a potent vasodilator, in the kininogen prior to the release of bradykinin.
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A; Residues: 450-452, Y, 454, Y, 456 <LIT>
A; Cross-references: UNIPARC: UPI000017429E
R; Straczek, J; Maachi, F; le Nguyen, D;
FEBS Lett. 373, 207-211, 1995
A; Title: Purification from human plasma of
A; Reference number: S68059; MUID: 96033974;
A; A; A; Ccession: S68059
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A;Reference number: 855239; MUID:95251593; P
A;Accession: 855239
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A;Cross-references: UNIDARC:UPI000002COF8
R;Lenarcic, B.; Krasovec, M.; Ritonja, A.; Olafsson, I.; Turk, V.
FEBS Lett. 280, 211-215, 1991
A;Title: Inactivation of human cystatin C and kininogen by human A;Reference number: S14303; MUID:91192133; PMID:2013314
A;Accession: S14447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Title: Structural features of plasma kinins and kininogens. A;Reference number: A91455; MUID:90255622; PMID:4952632 A;Contents: annotation; bradykinin
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A;Residues: 431-434 <STR>
A;Residues: 431-434 <STR>
A;Cross-references: UNIPARC:UPI000017429F
A;Cross-references: UNIPARC:UPI000017429F
T;Kltamura, N., Kitagawa, H.; Fukushima, D.; Takagaki, Y.; Miyata, T.; Nakanishi, J. Biol. Chem. 260, 8610-8617, 1985
J. Biol. Chem. 260, 8610-8617, 1985
A;Title: Structural organization of the human kininogen gene and a model for its of the formula organization of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a model for its of the human kininogen gene and a mode
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A; Residues: 264-359, N', 361-375 < LEN2>
A; Crose-references: UNIPARC: UPI000017429D
R; Little, 8.8.; Johnson, D.A.
B; Lottle, 8.8.; Johnson, D.A.
B; Little, 8.8.; Johnson, D.A.
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A;Residues: 380-389 <KAT1>
A;Cross-references: UNIPARC:UPI000002C0F8
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A;Experimental source: urine
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A;Residues: 1-19;189-192;310-314;381-389 <LENI>
A;Cross-references: UNIPARC:UPI00002CF4A; UNIPARC:UPI0000143284; UNIPARC:UPI0000174298
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Fed. Proc. 27, 52-57, 1968
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A, Residues: 380-389 < KAT3>
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ion and identification of hydroxyproline analogues of bradykinin
ber: A61495; MUID:88211869; PMID:3366244
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A;Cross-references: UNIPROT: D01046; UNIPARC: UPI000012DF24; GB: J00010; GB: V00426; NID: 916
C;Comment: The LMW kininogen precursor is produced from the same gene as the HMW form at C;Comment: Kininogen is a cysteine proceinase inhibitor, takes part in initiation of the C;Comment: Bradykinin, released from kininogen by kallikrein, is a potent vasodilator, a cyproline residue is present in the kininogen prior to the release of bradykinin.
C;Superfamily: kininogen; cystatin homology
C;Keywords: alternative splicing; blood coagulation; cysteine proteinase inhibitor; glyce;19-18/Domain: signal sequence #status predicted <SIG>F;1-18/Domain: signal sequence #status predicted <SIG>F;19-378/Product: LMW kininogen I #status predicted <HCH>F;19-378/Product: LMW kininogen I heavy chain #status predicted <HCH>F;19-378/Domain: cystatin homology <CY1>F;363-374/Domain: cystatin homology <CY2>F;363-374/Domain: cystatin homology <CY2>F;380-388/Product: LMW kininogen I heavy chain #status predicted <HBDY>F;380-388/Product: LMW kininogen I kallidin II) #status predicted <HBDY>F;380-388/Product: LMW kininogen I light chain #status experimental <LCH>F;379-318/Product: LMW kininogen I light chain #status experimental <LCH>F;19/Modified site: pyrrolidone carboxylic acid (Gln) (in mature form) #status predicted F;382-Modified site: Met-Lys (kallikrein) #status predicted F;388-389/Cleavage site: Met-Lys (kallikrein) #status predicted F;388-389/Cleavage site: Arg-Ser (kallikrein) #status predicted
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F;19-131/Domain: cystatin homology <CY1>
F;42-233/Domain: cystatin homology <CY2>
F;264-375/Domain: cystatin homology <CY3>
F;264-375/Domain: cystatin homology <CY3>
F;380-389/Product: lysyl-bradykinin (kallidin II) #status experimental <BDY>
F;380-389/Product: bradykinin (kallidin II) #status experimental <CCH>
F;380-389/Product: bradykinin (kallidin II) #status experimental <CCH>
F;380-389/Product: lowmolecular weight growth promoting factor #status experimental <F;421-510/Region: glycine/histidine/lysine-rich 30-residue repeats
F;431-434/Product: lowmolecular weight growth promoting factor #status experimental <F;19/Modified site: pyrrolidone carboxylic acid (Gln) (in mature form) #status experimental F;48/Binding site: carbohydrate (Asn) (covalent) #status absent F;48/Binding site: carbohydrate (Asn) (covalent) #status experimental F;383/Modified site: Met-Lys (kallikrein) #status experimental F;389-390/Cleavage site: Met-Lys (kallikrein) #status experimental F;389-390/Cleavage site: Arg-Ser (kallikrein) #status experimental F;401,531,542,546,557,571,593,628/Binding site: carbohydrate (Thr) (covalent) #status experimental
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N;Alternate names: alpha-2-thiol proteinase inhibitor; preprokininogen
N;Contains: bradykinin (kallidin); kininogen I; kininogen II; prokininogen
C;Species: Bos primigenius taurus (cattle)
C;Date: 14-Nov-1983 #sequence_revision 14-Nov-1983 #text_change 09-Jul-2004
C;Accession: A01283
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A;Accession: A01283
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A; Residues: 1-436 < NAW>
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGKKYFIDFVARETTCSKESNEBLTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP 122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98.5%; Scilarity 100.0%; F
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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prior to the release

of bradykinin

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Riningen, HMW I precursor - bovine
N;Alternate names: alpha-2-thiol proteinase inhibitor; preprokininogen
N;Contains: bradykinin (kallidin); kininogen I; kininogen II; prokininogen
C;Species: Bos primigenius teaurus (cattle)
C;Date: 14-Nov-1983 #sequence revision 14-Nov-1983 #text_change 09-Jul-2004
C;Accession: A01281; A91923; A91938; A29559
R;Kitamura, N.; Takagaki, Y.; Furuto, S.; Tanaka, T.; Nawa, H.; Nakanishi, S.
Nature 305, 545-549, 1983
A;Title: A single gene for bovine high molecular weight and low molecular weight kininog
A;Reference number: A93317; MUID:84014106; PMID:6571699
A;Accession: A01281
A;Molecule type: mMNA
A;Residues: 1-621 <KIT-
A;Cross-references: UNIPROT:P01044; UNIPARC:UPI000012DF1C; GB:V01491; GB:K01757; NID:949
A;Cross-references: UNIPARC:UPI00001742A1
A;Accession: A91923; MUID:70180420; PMID:4986212
A;Goss-references: UNIPARC:UPI00001742A1
R;Han, Y.N.; Komiya, M.; Iwanaga, S.; Suzuki, T.
J. Biochem. 67, 55-68, 1975
A;Title: Studies on the primary structure of bovine high-molecular-weight kininogen. Ami
A;Reference number: A91938; MUID:75170265; PMID:1169237
A;Accession: A91938
A;Molecule type: protein
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A; Residues: 458-498 cHAP.
A; CROSS-references: UNIPARC: UPI00001742A2
A; Cross-references: UNIPARC: UPI00001742A2
R; Sueyoshi, T.; Miyata, T.; Hashimoto, N.; Kato, H.; Hayash
T. Biol. Chem. 262, 2768-2779, 1987
J. Biol. Chem. 262, 2768-2779, 1987
A; Title: Bovine high molecular weight kininogen. The amino
A; Reference number: A92627; MUID:87137530; PMID:3546295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            δ
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                                                                                                                                                                                                     Seikagaku 56, 808, 1984
A;Title: Disulfide bonds in bovine
                                                                                                                                                                                                                                                                                                                    A;Title: The amino acid sequence of the light chain of human high-molecular-mass A;Reference number: A91153; MUID:86030270; PMID:4054110
                                                                                                                                                                                                                                                                                                                                                                                R;Lottspeich, F.; Kellermann, J.; Henschen, A.; Foertsch, Eur. J. Biochem. 152, 307-314, 1985
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;Note: article in Japanese; Comment: The HMW kininogen precursor is produced from the same gene as the LMW i; Comment: Kininogen is a cysteine proteinase inhibitor, takes part in initiation; Comment: The glycine/histidine/lysine-rich region of HMW kininogen light chain : Comment: Bradykinin, released from kininogen by kallikrein, is a potent vasodila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query
Best I
                                                                                                                                            Reference number: A94300 ;Contents: annotation; disulfide
                                                                                                                                                                                                                                                                                                                                                                                                                                      Molecule type: protein;Residues: 'Z',20-123,'I',125-127,'I',129-378;Residues: 'Z',20-123,'I',125-127,'I',129-378;Cross-references: UNIPARC:UPI00001742A3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                                                                                                                                                                                                                           annotation; bovine cleavage sites; T.; Miyata, T.; Kato, H.; Iwanaga,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4
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71.9%;
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Pred. No. 1.7e-37;
                                                                                                                                               ponds
                                                                                                                                                                                                        MMH
                                                                                                                                                                                                     kininogens
                                                                                                                                                                                                                                                                                               povine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        H.; Hayashida,
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                                                                                                                                                                                                                                                                                         carbohydrate binding sites
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                                                                                                                                                                                                                                                                                                                                                                                                              Muller-Esterl,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     H.; Miyata, T.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence,
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the LMW form as nitiation of the nit chain is import vasodilator, i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      positions
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F;19-137/Product: war homology <CY1>
F;19-130/Domain: cystatin homology <CY2>
F;141-252/Domain: cystatin homology <CY2>
F;261-372/Domain: cystatin homology <CY3>
F;261-372/Domain: cystatin homology <CY3>
F;377-386/Product: lysyl-bradykinin (kallidin II) #status predicted <KBDY>
F;378-386/Product: bradykinin (kallidin I) #status predicted <BDY>
F;378-386/Product: hradykinin (kallidin I) #status experimental <LCH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: mRNA
A;Residues: 1-434 <NAW>
A;Residues: 1-434 <NAW>
A;Residues: 1-634 <ANAWA
A;Cross references: UNIPROT:P01047; UNIPARC:UPI000012DF28; GB:V00427; GB:J00011; NID:94
A;Cross references: UNIPROT:P01047; UNIPARC:UPI000012DF28; GB:J00011; NID:94
A;Cross references: UNIPROT:P01047; UNIPARC:UPI000012DF28; GB:J00011; NID:94
A;Cross references: UNIPROT:P01047; UNIPARC:UPI000012DF28; GB:J00011; NID:94
A;Cross references: UNIPROT:P01047; UNIP
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kininogen, LMW II precursor - bovine
k;Alternate names: alpha-2-thiol proteinase inhibitor; preprokininogen
N;Contains: bradykinin (kallidin); kininogen I; kininogen II; prokininogen
C;Species: Bos primigenius taurus (cattle)
C;Date: 14-Nov-1983 #sequence_revision 14-Nov-1983 #text_change 09-Jul-2004
C;Date: 14-Nov-1983 #sequence_revision 14-Nov-1983 #text_change 09-Jul-2004
C;Accession: A01284
R;Nawa, H.; Kitamura, N.; Hirose, T.; Asai, M.; Inayama, S.; Nakanishi, S.
Proc. Natl. Acad. Sci. U.S.A. 80, 90-94, 1383
A;Title: Brinary structures of bovine liver low molecular weight kininogen ]
A;Reference number: A93984; MUID:83117859; PMID:6572010
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                                                                                                                                                                                                                                                                                                                                                C;Superfamily: kininogen; cystatin homology
C;Keywords: alternative splicing; blood coagulation; c;
C;Keywords: alternative splicing; blood coagulation; c;
F;1-18/Domain: signal sequence #status predicted <IGs
F;19-434/Product: LMW kininogen II #status predicted <I
F;19-377/Product: LMW kininogen I heavy chain #status |
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Pred. No. 2.5e-37;
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A;Cross-references: UNIPARC:UPI00001742A4
A;Note: 398-Pro, 401-Val, and 455-Lys were also found
R;Han, Y.N., Komiya, M.; Iwanaga, S.; Suzuki, T.
J. Biochem. 77, 55-68, 1975
A;Title: Studies on the primary structure of bovine high-molecular-weight kininogen.
A;Reference number: A91938; MUID:75170265; PMID:1169237
A;Accession: A91938
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N;Alternate names: alpha-2-thiol proteinase inhibitor; preprokininogen
N;Conteinas: bradykinin (kallidin); kininogen I; kininogen II; prokininogen
C;Species: Bos primigenius taurus (cattle)
C;Date: 14-Nov-1983 #sequence revision 14-Nov-1983 #text_change 09-Jul-2004
C;Date: 14-Nov-1983 #sequence revision 14-Nov-1983 #text_change 09-Jul-2004
C;Accession: A01282; A91923; F31941; A91938; B29559
R;Kitamura, N; Takagaki, Y.; Furuto, S.; Tanaka, T.; Nawa, H.; Nakanishi, instruct 305, 545-549, 1983
A;Molecule type: protein
A;Residues: 'Z', 20-104,'E', 106-256,'XX', 257-376
A;Cross-references: UNIPARC:UPI00001742A5
                                                                                                        A; Molecule type: protein
A; Residues: 456-496 < 4182>
A; Cross-references: UNIPARC; UPI00001742A2
A; Cross-references: UNIPARC; UPI00001742A2
R; Sueyoshi, T.; Miyata, T.; Hashimoto, N.; Kato, H.; Hayash
J. Biol. Chem. 262; 2768-2779; 1987
A; Title: Bovine high molecular weight kininogen. The amino
A; Reference number: A92627; MUID:87137530; PMID:3546295
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J. Biochem. 79, 1201-1222, 1976
A;Title: Primary structure of bovine plasma high-molecular-weight kininogen.
A;Reference number: A91941; MUID:76260155; PMID:956151
A;Accession: A91941
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A;Residues: 1-619 <KIT>
A;Residues: 1-619 <KIT>
A;Cross-references: UNIPROT:P01045; UNIPARC:UPI000012DF1F;
R;Kato, H.; Nagasawa, S.; Suzuki, T.
J. Biochem. 67, 313-323, 1970
J. Biochem. 67, 313-323, 1970
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A,Residues: 376-391 <KAT>
A,Cross-references: UNIPARC:UPI00001742A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PMID:4986212
                                                                                                                                                                                             H.; Hayashida,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cleavages
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                                                                                                                                                                                             H.; Miyata,
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F;441-222/Domain: Cystatin homology <CV2>
F;261-372/Domain: Cystatin homology <CV3>
F;261-372/Domain: Cystatin homology <CV3>
F;377-386/Product: lysyl-bradykinin (kallidin I) #status experimental <BDY>
F;377-386/Product: bradykinin (kallidin I) #status experimental <CV3>
F;387-619/Product: bradykinin (kallidin I) #status experimental <CV4>
F;387-619/Product: bradykinin (kallidin I) #status experimental <CV4>
F;387-619/Product: bradykinin (kallidin I) #status experimental <CV4>
F;387-619/Product: bradykinin (kallidin #status experimental <CV4>
F;387-619/Product: bradykinin (kallidin #status experimental <CV4+
F;387-619/Product: bradykinin (kallidin #status experimental <CV4+
F;387-619/Product: bradykinin (kallidin #status experimental #status experimental #status experimental #status experimental F;376-317/Cleavage site: carbohydrate (Asn) (covalent) #status experimental F;386-387/Cleavage site: Arg-Ser (kallikrein) #status experimental F;396,400,404,510/Binding site: carbohydrate (Ser) (covalent) #status experimental F;396,400,404,510/Binding site: carbohydrate (Ser) (covalent) #status experimental F;397,398,518,522,534,546,551,568/Binding site: carbohydrate (Ser) (covalent) #status experimental #status expe
                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 7
A28055
K-kininogen,
A;Reference number: A92496;
A;Accession: A28055
A;Molecule type: mRNA
A;Residues: 1-433 <FUR>
                                                                                                                                                                                                                                                                                                K-kininogen, LMW I precursor - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 20-Jun-1989 #sequence_revision 20-Jun-1989 #text_change
C;Accession: A28055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Note: article in Japanese C;Comment: The HAW kininogen precursor is produced from the same gene as the LAW form as C;Comment: Kininogen is a cysteine proteinase inhibitor, takes part in initiation of the C;Comment: Kininogen is a cysteine/lysine-rich region of HAW kininogen light chain is import;C;Comment: Bradykinin, released from kininogen by kallikrein, is a potent vasodilator, yxproline residue is present in the kininogen prior to the release of bradykinin. C;Superfamily; kininogen; cystatin homology.
                                                                                                                                                                                                                                    R;Furuto-Kato, S.; Matsumoto, A.; Kitamura, J. Biol. Chem. 260, 12054-12059, 1985
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F;19-619/Product: HMW kininogen II #status predicted <MAT-
F;19-376/Product: HMW kininogen II heavy chain #status exp-
F;19-130/Domain: cystatin homology <CY1-
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A;Title: The amino acid sequence of the light chain of human high-molecular-mass
A;Reference number: A91153; MUID:86030270; PMID:4054110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Contents: annotation; disulfide bonds
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A;Title: Disulfide bonds in bovine HMW kininogens.
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Best Local :
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68.9%;
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Pred. No. 9.1e-35;
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                                                                                                                                                                                                                                                                            N.; Nakanishi,
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UNIPROT: P08934; UNIPARC: UPI0000177A66

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C; Superfamily: kininogen; cystatin homolog C; Keywords: alternative splicing F;1-18/Domain: signal sequence #status preF;19-433/Product: K-kininogen, LMW I #stat F;19-131/Domain: cystatin homology <CY1>F;142-253/Domain: cystatin homology <CY2>F;264-375/Domain: cystatin homology <CY3>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Title: Differing expression patterns and evolution of the rat kininogen gene family. A; Reference number: A92625; MUID:87137443; PMID:3029068
A; Accession: A25486
A; Molecule type: mRNA
A; Residues: 1-639 <KIT>
A; Residues: 1-639 <KIT>
A; Cross-references: UNIPROT:P08934; UNIPARC:UPI000012DF1B
A; Otto: the authors translated the codon CAA for residue 347 as Asn C; Superfamily: kininogen; cystatin homology
C; Keywords: alternative splicing
F; 19-639/Product: kininogen, HAW I #status predicted <NAT>
F; 19-131/Domain: cystatin homology <CY1>
F; 19-131/Domain: cystatin homology <CY1>
F; 19-131/Domain: cystatin homology <CY1>
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C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 08-Mar-1989 #sequence_revision 08-Mar-1989 #text_change 09-Jul-2004
C;Date: 08-Mar-1989 #sequence_revision 08-Mar-1989 #text_change 09-Jul-2004
C;Date: 08-Mar-1989 #text_change 09-Jul-2004
C;Accession: A25486
R;Kitagswa, H.; Kitamura, N.; Hayashida, H.; Miyata, T.; Nakanishi, S.
J. Biol. Chem. 262, 2190-2198, 1987
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A25486
kininogen, HMW I precursor -
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                major acute phase alpha-1 protein (version 2) - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 19-May-1989 #sequence_revision 19-May-1989 #text_change 09-Jul-2004
C;Accession: A23897; B23897
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R; Anderson,
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Matches 80; Conserv
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 426; DB 2;
Pred. No. 1.8e-32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 426; DB 2;
Pred. No. 1.2e-32;
                                                                                                                                                                                                                                                                                                                                                                                                              <u>:-</u>
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A;Cross-references: UNIPROT:P08932; UNIPARC:UPI000012DF30 R;Kitagawa, H.; Kitamura, N.; Hayashida, H.; Miyata, T.; N.; Biol. Chem. 262, 2190-2198, 1987
A;Title: Differing expression patterns and evolution of the A;Reference number: A92625; MUID:87137443; PMID:3029068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C;Species: Rattus norvegicus (Norway rat)
C;Date: 20-Jun-1989 #sequence revision 20-Jun-1989 #text_change 09-Jul-2004
C;Accession: B28055; E25486; B28526; C28526
R;Furuto-Kato, S.; Matsumoto, A.; Kitamura, N.; Nakanishi, S.
J. Biol. Chem. 260, 12054-12059, 1985
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Note: the authors translated the codon C;Superfamily: kininogen; cystatin homolc F;19-130/Domain: cystatin homology <CY2-F;141-252/Domain: cystatin homology <CY2-
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A; Residues: 5-430 <AND2>
A; Cross-references: UNIPARC: UPI0000170A0C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: UNIPROT:P01048; A;Accession: B23897
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A;Title: The relationship between rat major acute phase A;Reference number: A23897; MUID:86008266; PMID:2413019
A;Accession: A23897
A;Molecule type: protein
A;Residues: 1-14 <AND1>
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                     C;Superfamily: kininogen; cystatin homology C;Keywords: glycoprotein; pyroglutamic acid F;1-18/Domain: signal sequence #status predi
                                                                                                 A; Molecule type: protein
A; Residues: 'E', 20-48; 376-388, 'R', 390-419, 'ER', 422-430
A; Cross-references: UNIPARC: UPI0000177A67
                                                                                                                                                                                                                                                                                                   A;Cross-references: UNIPARC:UPI000000380
R;Enjyoji, K.; Kato, H.; Hayashi, I.; Oh-ishi, S.; Iwanaga, S.
J. Biol. Chem. 263, 973-979, 1988
A;Title: Purification and characterization of rat T-kininogens
A;Reference number: A92729; MUID:88087226; PMID:3121623
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F;263-374/Domain: cystatin homology <CY3>
                                                                                                                                                                                                    A;Molecule type: protein
A;Residues: 'E', 20-25, 'MD', 28-48,376-430 <ENJ>
A;Cross-references: UNIPARC:UPI0000177A67
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A; Residues: 1-430 < FUR>
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A;Accession: B28055
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                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Molecule type: DNA
A;Residues: 375-430 <KIT>
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F;19-430/Product:
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Best Local
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  T-kininogen,
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     EME
                            #status predicted <SIG>
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Pred. No. 4.7e-31;
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     #status
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                evolution of the PMID:3029068
     experimental <MAT>
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R;Cole, T.; Inglis, A.S.; Roxburgh, C.M.; Howlett, G.J.; Schreiber, FBBS Lett. 182, 57-61, 1985
"A;Title: Major acute phase alphal-protein of the rat is homologous 'A;Reference number: A01285; MUID:85127561; PMID:2578992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          r.
C;Superfamily: kininogen; cystatin homology
C;Keywords: bradykinin; cysteine proteinase
P;1-11/Domain: signal sequence (fragment) #c
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F;263-274/Domain: cystatin homology <CY3>
F;263-274/Domain: cystatin homology <CY3>
F;19/Modified site: pyrcolidone carboxylic acid (Gln) (in mature form) #status experimen F;82,126,168,204,326/Binding site: carbohydrate (Asn) (covalent) #status predicted F;82,126,168,204,326/Binding site: carbohydrate (Asn) (covalent) #status predicted F;83-94,107-125,141-144,205-217,228-247,263-266,327-339,350-369/Disulfide bonds: #status
T-kininogen
N,Alternate
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À;Molecule type: mRNA
A;Residues: 1-423 <COL>
À;Cross-references: UNII
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N;Contains: bradykinin
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P,141-252/Domain: cystatir
F,263-374/Domain: cystatir
                                              KGRTT1
                                                                   RESULT 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  134-245/Domain: cystatin homology <CY2>
;256-367/Domain: cystatin homology <CY3>
;371-379/Product: bradykinin #status predicted <BDY>
;12/Modified site: pyrrolidone carboxylic acid (Gln)
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Best Local
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Best Local
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I precursor names: 73K p
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                                                                                                                                                                                                                                                                      AGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQP 122
                                                                                                                                                                                                                                                                                                                GDDLFELLPKNCRGCPREIPVDSPELKEALGHSIARLNAQHNHIFYFKIDTVKKATSQVV
                                                                                                                                                                                                                                                                                                                                                               GKDFVQPPTKICVGCPRDIPTNSPELEETLTHTITKLNAENNATFYFKIDNVKKARVQVV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LGM 125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GDDLFSLLPKKCFGCPKNIPVDSPELKEALGHSIAQLNAQHNHLFYFKIDTVKKATSQVV
                                                                                                                                                                                    LGM 125
                                                                                                                                                                                                                           AGVIYVIEFIARETNOSKOSKTELTADOETKHLGOSLNONANVYMRÞWENKVVÞTVROOA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LDM 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              plasma glycoprotein inhibits cysteine proteinases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                           60.0%;
                       rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         homology <CYl>
  EME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 409; DB 2; Length 430 Pred. No. 4.7e-31; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 402; DB 1;
Pred. No. 2.1e-30;
  kininogen
                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27-Nov-1985 #text_change 12-Apr-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rat (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 423
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (in mature form) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      homologous to
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A;Molecule type: mRNA
A;Residues: 330-420,'R',422-429,'P' <KAN>
A;Residues: 330-420,'R',422-429,'P' <KAN>
A;Residues: 330-420,'R',422-429,'P' <KAN>
A;Residues: 330-420,'R',422-429,'P' <KAN>
A;References: UNIPARC: UPI00001742A9
R;Anderson, K.P.; Croyle, M.L.; Lingrel, J.B.
Gene 81, 119-128, 1989
R;Anderson, K.P.; Croyle, M.L.; Lingrel, J.B.
Gene 81, 119-128, 1989
A;Reference number: JQ0027; MUID:90034172; PMID:2806908
A;Accession: JQ0027
A;Accession: JQ0027
A;Accession: JQ0027
A;Residues: 1-60,'E',62-113,'R',115-165,'F',167-178,'TKI',182-211,'F',213-256,'S',258-31
A;Residues: 1-60,'E',62-113,'R',115-165,'F',167-178,'TKI',182-211,'F',213-256,'S',258-31
A;Cross-references: UNIPARC:UPI00001742AA
A;Experimental source: strain Sprague-Dawley
R;Kageyama, R.; Kitamura, N.; Ohkubo, H.; Nakanishi, S.
J. Biol. Chem. 262, 2345-2351, 1987
A;Title: Differing utilization of homologous transcription initiation sites of rat K and A;Reference number: A25488; MUID:87137465; PMID:3818598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: UNIPARC:UPI000000041B; GB:M14356; NID:g205090; PIDN:AAA41492.1; R;EnjyOji, K.; Kato, H.; Hayashi, I.; Oh-ishi, S.; Iwanaga, S. J. Biol. Chem. 263, 965-972, 1986 J. Biol. Chem. 263, 965-972, 1986 A;Title: Purification and characterization of two kinds of low molecular weight kin: A;Reference number: A28525; MUID:88087225; PMID:3335530
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 17-Mar-1987 #sequence revision 17-Mar-1987 #text_change 31-Dec-2004
C;Date: 10-Mar-1987 #sequence revision 17-Mar-1987 #text_change 31-Dec-2004
C;Accession: A01286; D26486; A28526; PL0193; JQ0027; B25488; A28525; S68036
R;Furuto-Kato, S.; Matsumoto, A.; Kitamura, N.; Nakanishi, S.
J. Biol. Chem. 260, 12054-12059, 1985
                                                                                                                                                                                                                                                                                                                                                 A;Molecule type: protein
A;Residues: 376.430 <ENDIANC:UPI00001742A8
A;Cross-references: UNIPARC:UPI00001742A8
A;Cross-references: UNIPARC:UPI00001742A8
A;Sierra, F.; Walter, R.; Vautravers, P.; Guigoz, Y.
A;Ch. Biochem. Biophys. 322, 333-338, 1995
A;Tile: Identification of several isoforms of T-kininogen
A;Reference number: S68034; MUID:96032652; PMID:7574705
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R;Kanda, S.; Sugiyama, K.; Takahashi, M.; Shumiya, S.; Tomino, S.; Nagase, S.
Jpn. J. Cancer Res. 81, 63-68, 1990
A;Title: Identification of a protein increasing in serum of Nagase analbuminemic rats
A;Reference number: PL0193; MUID:90216390; PMID:2108948
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R;EnjyOji, K.; Kato, H.; Hayashi, I.; Oh-ishi, S.; Iwanaga, S.
J. Biol. Chem. 263, 973-979, 1988
A;Title: Purification and characterization of rat T-kininogens
A;Reference number: A92729; MUID:88087226; PMID:3121623
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A;ROBIGURES: 1-430 <FUR>
A;ROBIGURES: 1-430 <FUR>
A;Cross-references: UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references:UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references:UNIPROT:P01048; UNIPROT:Q63283; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references:UNIPROT:P01048; UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references:UNIPROT:Q63283; UNIPARC:UPI0000163B79;
A;Cross-references:Q64285; UNIPARC:UPI0000163B79;
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A;Cross-references:Q64285; UNIPARC:UPI0000163B79;
A;Cross-references:Q64285; UNIPARC:UPI0000163B79;
A;Cross-references:Q64285; UNIPARC:UPI0000163B79;
A;Cross-references:Q64285; UNIPARC:UPI0000163B79;
A;
A;Cross-references: UNIPARC:UPI00001742AB
A;Experimental source: clone pSG17
C;Comment: At least three types of LMW ki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Accession: B25488
A;Status: preliminary
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A;Title: Differing expression patterns and A;Reference number: A92625; MUID:87137443;
                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 340-430 <SIE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-48 < KAG>
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A; Residues: 'E', 20-48; 376-430 < ENJ>
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A; Residues: 375-430 < KIT>
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of LMW kininogen

precursors are

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rat

plasma,

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ceding bradykinin.

C;Comment: T-kiningens contain T-kinin (I-S-bradykinin), a novel kinin isolated C;Comment: T-kiningens contain T-kinin (I-S-bradykinin), a novel kinin isolated d of an Arg or Lys, it is probably not released from its precursor by either tiss C;Comment: The T-kiningens are produced in response to an inflammatory stimulant C;Genetics:

C;Genetics:
C;Genetics:
A;Introns: 65/3; 102/3; 130/1; 187/3; 223/2; 252/1; 309/3; 345/3; 374/3; 398/3
C;Superfamily: cystatin homology
C;Keywords: acute phase; bradykinin; cysteine proteinase inhibitor; duplication; C;Keywords: acute phase; bradykinin; cysteine proteinase inhibitor; duplication; F;11-18/Domain: signal sequence #scatus predicted <SIG>
F;19-430/Product: T-kininogen I #scatus experimental <MAT>
F;19-130/Domain: Cystatin homology <CYI>
CYCII-CSCATIN homology <CYI>
CYCII-CSCATIN homology <CYI>
CYCII-CSCATIN homology <CYII-CSCATIN homology <CYII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Cross-references: UNIPARC:UPI0000173259
C;Superfamily: cystatin; cystatin homology
C;Keywords: colostrum; cysteine proteinase
F;2-112/Domain: cystatin homology <CYS-
F;48-52/Region: inhibitory #status predicte
F;66-76,90-110/Disulfide bonds: #status pre
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F;263-374/Domain: cystatin homology <CY3>
F;378-386/Product: bradykinin #status predicted <BDY>
F;378-386/Product: bradykinin #status cystatic acid (Gln) (in mature form) #status experimen F;19/Modified site: pyrrolidone carboxylic acid (Gln) (covalent) #status predicted F;82,126,168,204,326/Binding site: carbohydrate (Asn) (covalent) #status predicted F;83-94,107-125,141-144,205-217,228-247,263-266,327-339,350-369/Disulfide bonds: #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N;Alternate names: thiol proteinase inhibitor
C;Species: Bos primigenius taurus (cattle)
C;Date: 28-Feb-1986 #sequence_revision 28-Feb-1986 #text_change 06-Dec-1996
C;Accession: A01271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 გ
RESULT 14
$68035
T-kininogen (clone pSG17) - rat (frac
C;Species: Rattus norvegicus (Norway
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Title: Complete amino acid sequence of bovine colostrum low-M-r cysteine proteinase A;Reference number: A01271; MUID:85231205; PMID:3891407 A;Accession: A01271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Accession: A01271
R;Hirado, M.; Tsunasawa,
FEBS Lett. 186, 41-45, 1
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Best Local S
Matches 76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ,2-112/Domain: cystatin homology <CYS>
,48-52/Region: inhibitory #status predicted
,66-76,90-110/Disulfide bonds: #status predicted
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Best Local :
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ilarity 30.6%;
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       rat (fragment)
s (Norway rat)
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Pred. No. 7.7e-06;
2; Mismatches 31
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C;Date: 12-Feb-1998 #sequence_revision 20-Feb-1998 #text_change 31-Dec-2004 C;Accession: S68035  
C;Accession: S68035  
R;Sierra, F.; Walter, R.; Vautravers, P.; Guigoz, Y. Arch. Biochem. Blophys. 322, 333-338, 1995  
A;Fitle: Identification of several isoforms of T-kininogen expressed in the A;Reference number: S68034; MUID:96032652; PMID:7574705  
A;Accession: S68035  
A;Molecule type: mRNA A;Residues: 1-91 <SIE> A;Cross-references: UNIPROT:P70517; UNIPROT:Q63581; UNIPARC:UPI0000177A60  
A;Cross-references: UNIPROT:P70517; UNIPROT:Q63581; UNIPARC:UPI0000177A60
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                                                                                                                                                                                                                                                                                                                     Arch. Biochem. Biophys. 322, 333-338, 1995
A;Title: Identification of several isoforms of T-kininogen
A;Reference number: S68034; MUID:96032652; PMID:7574705
                                                                                                                                                                                                                                                                                                                                                                                                      T-kininogen (clone pSG22) - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 12-Feb-1998 #sequence_revision 20-Feb-1998 #text_change 31-Dec-2004
C;Accession: S68034
                                                                                                                                                                   A;Experimental source: liver
C;Superfamily: cystatin homology
C;Keywords: alternative splicing; cysteine proteinase inhibitor;
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A;Residues: 1-91 <SIE>
A;Cross-references: UNIPROT:P70517; UNIPROT:Q63581; UNIPARC:UPI0000177A5F
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                        91 ETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQPLGM 125
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1 ETKHLGQSLNCNANVYMRPWENKVVPTVRCQALDM 35
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 UniProt_05.80:*
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Q53EO HUMAN
Q6PAU9 HUMAN
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KNG2 BOVIN
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Q6S971 MOUSE
Q5MBAO RAT
KNG1 RĀT
Q6S970 MOUSE
Q6S913 MOUSE
Q58913 MOUSE
Q5MB4 RAT
Q7XRP6 PIG
KNT1 RĀT
Q7XRP6 PIG
KNT1 RĀT
Q5PQ11 RAT
Q5PQ11 RAT
Q5PQ11 RAT
Q5PQ12 TETNG
CYTF MOUSE
Q5PU15 MOUSE
Q5PU16 XENTR
Q7XFF HŪMAN
Q5XJ76 BRARE
Q5MB73 XENTR
Q4SPM6 TETNG
Q7SYM2 XENLA
Q7ZY91 XENLA
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               Q5m894
P0197p6
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P0197p4
Q5pqu1
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Q64std5
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Q9qw15
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LOCAL SIMILATITY 100.0%; PFEG. NO. BE 123; CONSERVATIVE 0; MISMACC GROPPOPPKICVGCPRDIPTNSPELEETL GROPPOPPKICVGCPRDIPTNSPELEETL 63 AGKKYFIDFVARETTCSKESNBELTESCET	Heruction and characterization of deniched cDNA library."; ne 200:149-156(1997). CLEOTIDE SEQUENCE. SSUB=Kidney; coki Y., Toyoda A., Takeda T., Sakabonitted (ApR-2005) to the EMBI/GenBBBL, AK223589; BAD97309.1; -; mRNA. N TER 1 N TER 1 OBSASS Match 98 5%: Score 660	COMO.	ALIGNMENT HUMAN HUMAN J-SBQO (Tremblrel. 31, Created) J-SEP-2005 (Tremblrel. 31, Last sequence) J-SEP-2005 (Tremblrel. 31, Last annotate) (Aninogen 1 variant (Fragment). John Seplens (Human). Sukaryota; Metazoa; Chordata; Craniata; Jammalia; Eutheria; Evarchontoglires; Pr	2 152.5 22.8 586 2 Q5X 3 148.5 22.2 318 2 Q4S 4 138.5 20.7 149 2 Q6T 5 138.5 20.7 149 2 Q5T 6 138.5 20.7 149 2 Q5T 7 136 20.3 148 1 CYT 9 133.5 19.9 146 1 CYT 0 132.5 19.8 146 1 CYT 131 19.6 454 2 Q6T 2 131 19.6 454 2 Q6T 3 130 19.4 127 1 CYT 4 130 19.4 140 2 Q6T 4 130 19.4 140 2 Q6T 5 129.5 19.3 140 1 CYT
53; Indels LNAENNATFYFKI LNAENNATFYFWE LDCNAEVYVVPWE LDCNAEVYVVPWE	anaka A., Yokoyama S.; databases. E9 CRC64; Iength 427:	the cap structure of A., Sugano S.; A., Sugano S.;	'7 AA. '9 e update) ion update) Vertebrata; Euteleostomi; imates; Catarrhini; Hominidae;	Q5xka7 xenopus lae Q4smm7 tetraodon n Q15828 homo sapien Q6ibd2 homo sapien Q540n7 homo sapien Q540n7 homo sapien p01035 bos taurus Q54826 bos taurus Q54826 bos taurus Q19092 macaca mula Q19093 saimiri sci Q6f6td bitis gabon Q66107 xenopus lae P14841 rattus norv Q5m968 rattus norv P21460 mus musculu

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                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 123
1.HUMAN S
KNG1 HUMAN S
P01042; P01043;
21-JUL-1986 (Rel
01-FEB-1996 (Rel
                                                                                                                                                                                                                                                                                                                                                                                                                                              Strausberg R.;
Submitted (OCT-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; BC060039, AAH60039.1; -; mRNA.
GO; GO:0004869; F:cysteine protease inhibitor activity;
InterPro; IPR000010; Prot inh_cystat.
                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00031; Cystatin; 3.
SMART; SM00043; CY; 3.
PROSITE; PS00287; CYSTATIN;
SEQUENCE 427 AA; 47901 M
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Eukaryota; Metazoa;
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                                                   STANDARD;
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 33,
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Last sequence update)
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Pred. No. 2.2e-53;
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NUCLEOTIDE SEQUENCE [MRNA] (ISOFORM LMW).

MEDLINE=85122621, PubMed=6441591;
Ohkubo I., Kurachi K., Takasawa T., Shiokawa H., Sar
"Isolation of a human cDNA for alpha 2-thiol proteir
its identity with low molecular weight kininogen.";
Biochemistry 23:5691-5697(1984).
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Eur.
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"SeattleSNPs. NHLBI HL66682 program for genomic applications,";
FHCRC, Seattle, WA (URL: http://pga.gs.washington.edu).";
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Kininogen-1 precursor (Alpha-2-thiol proteinase
Kininogen-1 heavy chain; Bradykinin (Kallidin I)
(Kallidin II); Kininogen-1 light chain; Low mole
promoting factor].
Name=KNG1; Synonyms=BDK, KNG;
Nabet P., Belleville ...
"Purification from human plasma
"nsulin-like growth factor-I act
PEBS Lett. 373:207-211(1995).
                                                                                                                                                                                                                                                                                                                                                             ASN-205, AND ABSENCE OF CARBOHYDRATE AT ASN-48.

MEDILYBE-86108361; PubMed-3484703;

Kellermann J., Lottspeisch F., Henschen A., Muller-Ester

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evidence for its evolution by gene triplication.";

Eur. J. Biochem. 154:471-478(1986).
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                                                                                                                                                                                                                     PROTEIN SEQUENCE OF 380-389, AND HYDROXYLATION SITE PRO-383. MEDLINE=88211869; PLDMed=3366244; DOI=10.1016/0014-5793(88)80427-7; MAEDLINE, MAEDLIN
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MEDLINE=86030270; PubMed=405
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                                                                                                         Straczek J., Maachi F.,
                                                                                                                                MEDLINE=96033974;
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J. Biochem.
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27:52-57(1968).
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                                                                                                                                OF 431-434, AND I
; PubMed=7589467;
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oglires; Primates; Catarrhini; Hominidae;
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                                            of a tetrapeptide that potentiates tivity in chick embryo cartilage.";
                                                                                                         MASS SPECTROMETRY.
, DOI=10.1016/0014-5793
n D., Becchi M., Heulin
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structures
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Protections 4:45.465(2004).

C -i- FUNCTION: (1) Kininogens are inhibitors of thiol proteases; (2)

C -i- FUNCTION: (1) Kininogens are inhibitors of thiol proteases; (2)

C -i- FUNCTION: (1) Kininogens are inhibitors of thiol proteases; (2)

C -i- FUNCTION: (2) Kininogens inhibits the coagulation by

the liping to position optimally prekallikrein and factor XI next to

factor XII; (3) HMW-kininogen inhibits the thrombin-and plasmin-

c induced aggregation of thrombocytes; (4) the active peptide

contraction, that is released from HMW-kininogen shows a variety of

c physiological effects: (4A) influence in smooth muscle

contraction, (4B) induction of hypotension, (4C) natriuresis and

c diuresis, (4D) decrease in blood glucose level, (4E) it is a

mediator of inflammation and causes (4E1) increase in vascular

permeability, (4E2) stimulation of nociceptors (4E3) release of

other mediators of inflammation (e.g. prostaglandins), (4F) it has

c chardioprotective effect (directly via bradykinin action,

indirectly via endothelium-derived relaxing factor action); (5)

LMW-kininogen is in contrast to HMW-kininogen not involved in blood
     EMBL; K02566;
EMBL; M11437;
EMBL; M11438;
                                                                                                                                                                                                                                                                                                                This Swiss-Prot entry is copyright. It is produced through a continuous the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict
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MEDLINE-22660472; PubMed=12754519; DOI=10.1038/nbt827;
Zhang H., Li X.-J., Martin D.B., Jebersold R.;

"Identification and quantification of N-linked glycoprohydrazide chemistry, stable isotope labeling and mass;
hydrazide chemistry, stable isotope labeling and mass;
Nat. Biotechnol. 21:660-666(2003).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CARBOHYDRATE-LINKAGE SITES ASN-169 AND ASN-294, PubMed=14760718; DOI=10.1002/pmic.200300556; Bunkenborg J., Pilch B.J., Podtelejnikov A.V.,
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SUBCELULIAR LOCATION: Secreted; extracellular.
ALTERNATIVE PRODUCTS:
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SSUE SPECIFICITY: Plasma.
W: Bradykinin is released from kininogen
W: Hydroxylation of Pro-383 occurs prior
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56:808-808(1984).
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DR EMBL; M11522; AABS955.
DR EMBL; M11522; AABS9551.1;
DR EMBL; M11523; AABS9551.1;
DR EMBL; M11524; AABS9551.1;
DR EMBL; M11526; AABS9551.1;
DR EMBL; M11526; AABS9551.1; JOL.
DR EMBL; M11527; AABS9551.1; JOL.
DR EMBL; M1527; AABS9551.1; JOL.
DR EMBL; M1528; AABS9551.1; JOL.
DR EMBL; M1528; AABS9551.1; JOL.
DR EMBL; A01240; KGHUHI.
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Best Local Sim
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HGNC; HGNC:6383; KNGI.
Reactome; P01042; -.
MIM; 228950; -.
GO:0005576; C:extrac
GO; GO:0008201; F:recept
GO; GO:0008270; F:recept
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GO; GO:0006954; P:indirac
GO; GO:0030147; P:natrai
GO; GO:0030147; P:negati
GO; GO:0043065; P:positi
GO; GO:0043065; P:positi
GO; GO:0006939; P:smooth
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Alternative splicing; Blood coagulation; Bradykinin;
Direct protein sequencing; Glycoprotein, Hydroxylation;
Inflammatory response; Plasma; Polymorphism; Protease i
Pyrrolidone carboxylic acid; Repeat; Signal; Thiol prot
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SIGNAL 1 18
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GO: GO:0005576; C:extracellular region; NAS.
GO; GO:0005201; F:heparin binding; NAS.
GO; GO:0005102; F:receptor binding; IPI.
GO; GO:0005102; F:receptor binding; NAS.
GO; GO:0005102; F:receptor binding; NAS.
GO; GO:000146; P:diuresis; TAS.
GO; GO:000146; P:inflammatory response; TAS.
GO; GO:000594; P:inflammatory response; TAS.
GO; GO:00030147; P:negative regulation of blood
GO; GO:0007162; P:negative regulation of cell a
GO; GO:00043065; P:positive regulation of apopto
GO; GO:0006939; P:smooth muscle contraction; TA
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InterPro; IPR000010; Prot inh c
Pfam; PF00031; Cystatin; 3.
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ilarity 100.0%;
Conservative
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f cell adhesion;
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KNG1_BOVIN ST/ P01044; P01046; 21-JUL-1986 (Rel.

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RP PROTEIN SEQUENCE OF 458-498.

RX MEDLINE-75170265; PubMed=1169237;
RX MEDLINE-75170265; PubMed=1169237;
RA Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
RA Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
RA Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
RA Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
RA Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
RI Stininogen. Amino acid sequence of a fragment ('histidine-rich RT kininogen. Amino acid sequence of a fragment ('histidine-rich RT RT peptide') released by plasma kallikrein.";
RI J. Biochem. 77:55-68(1975).
CC .-- PUNCTION: (1) Kininogens are inhibitors of thiol proteases; (2)
CC .-- PUNCTION: (1) Kininogens are inhibitors of thiol proteases; (2)
CC .-- PUNCTION: (1) Kininogens are inhibitors of thiol proteases; (2)
CC .-- PUNCTION: (1) HWW-kininogen inhibits the thrombin-and plasmin-
CC factor XII; (3) HWW-kininogen inhibits the active peptide
CC factor XII; (3) HWW-kininogen inhibits the active peptide
CC physiological effects (4A) influence in smooth muscle
CC physiological effects (4A) influence in smooth muscle
CC mediator of inflammation of hypotension, (4C) natriuresis and
CC mediator of inflammation and causes (EI) increase in vascular
CC permeability, (4E2) stimulation of nociceptors (4E3) release of
CC cother mediators of inflammation (e.g. prostaglandins), (4F) it has
CC indirectly via endocthelium-derived relaxing factor action); (5)
CC indirectly via endocthelium-derived relaxing factor action); (5)
CC kininogen is in contrast to HWW-kininogen not involved in blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kato H., Nagasawa S., Suzuki T.;
"Studies on the structure of bovine kininogen: cleavages bonds and of methionyl bonds in kininogen-II.";
J. Blochem. 67:313-323(1970).
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13-SEP-2005 (Rel. 48, Last annotation update)
Kininogen-1 precursor (Kininogen I) (Thiol proteinase inhibitor)
[Contains: Kininogen-1 heavy chain; Bradykinin (Kallidin I); Lyst
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sueyoshi T., Miyata T., Hashimoto N., Kato H., Hayashida H., Miyata T., Iwanaga S.; "Bovine high molecular weight kininogen. The amino acid sequ positions of carbohydrate chains and disulfide bridges in the chain portion.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-84014106; PubMed=6571699;
Kitamura N., Takagaki Y., Puruto S., Tana
"A single gene for bovine high molecular
weight kininogens.";
Nature and severe (MRNA) (ISOFORM HMW).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE-83117859, PubMed-6572010;
Nawa H., Kitamura N., Hirose T., Asai M., In
"Primary structures of bovine liver low mole
precursors and their two manas.";
Proc. Natl. Acad. Sci. U.S.A. 80:90-94(1983)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nature
[2]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Name=KNG1;
                                                                         <del>-</del> <del>-</del> <del>-</del>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ASN-168; ASN-169; ASN-197 AND ASN-204.
MEDLINE=87137530; PubMed=3546295;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROTEIN SEQUENCE OF 19-378,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE [MRNA]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Biol.
                                 - SUBCELLULAR LOCATIO
- ALTERNATIVE PRODUCT
Event=Alternative E
Name=HMW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         305:545-549(1983).
                   IsoId=P01044-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ta; Metazoa; Chorda
a; Eutheria; Laura
Bovidae; Bovinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Bovine).
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                                                                           LOCATION:
PRODUCTS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chordata; Crani.
Laurasiatheria;
                                                     splicing;
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                                                                                             Secreted;
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heria; Cetartiodactyla; Ruminantia;
                                                         Named
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , Tanaka T., I
cular weight
                                                                                               extracellular.
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v molecular
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nawa H., Nakanishi S.; and low molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              S., Nakanishi S.; weight kininogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SITES ASN-87;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of disulfide
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Alternative splicing; Blood coagulation; Bradykinin; birect protein sequencing; Glycoprotein; Inflammatory plasma; Protease inhibitor; Pyrrolidone carboxylic aci. Signal; Thiol protease inhibitor; Vasoactive; Vasodila: SIGNAL 1 18 Probable. CHAIN 19 621 Kininogen-1. Kininogen-1 heavy chain.
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CARBOHYD
CARBOHYD
CARBOHYD
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PEPTIDE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This Swiss-Prot entry is copyright. It is produced through a collabor between the Swiss Institute of Bioinformatics and the EMBL outstate European Bioinformatics Institute. There are no restrictions cuse as long as its content is in no way modified and this statement
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DISULFID
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CARBOHYD
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CARBOHYD
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PIR; A01283; KGBOLJ
   CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SMART; SM00043; CY;
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InterPro; IPR000010; Prot_inh_cystat.
Pfam; PF00031; Cystatin; 3.
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                                                                                                                                                                                  DISULFID
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                                   VARSPLIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE SPECIFICITY: Plasma.

PTM: Bradykinin is released from kininogen by SIMILARITY: Contains 3 cystatin-like domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IsoId=P01044-2; Sequence=VSP_013562, VSP_013563;
SSUB SPECIFICITY: Plasma
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295
621
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204
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520
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                                   621
      68890
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                                                                                                                                                                                         Cystatin-like 1.
Cystatin-like 3.
Cystatin-like 3.
Cystatin-like 3.
Cleavage (by kallikrein)
Pyrrolidone carboxylic a
N-linked (GlCNAC...)
N-linked (GlCNAC...)
N-linked (GlCNAC...)
N-linked (GlCNAC...)
N-linked (GlCNAC...)
O-linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mRNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lysyl-bradykinin.
Bradykinin.
Kininogen-1 light
                       VSLPHSAMSPVQDEERDSGKEQGPTHGHGWDHGKQI
HVKSCEYKGRPQEAGAEPAPQGEVSLPAESPQLAR
isoform LMW).
/FTId=VSP 013562.
Missing (In isoform LMW).
/FTId=VSP 013563.
                                                                                                                                                                                    Interchain
       D16850BEFE3C55CD CRC64;
                 VSP 013563.
(in Ref. 2
                                                                                                                                                                                                                                                                                                                                                                                    (by kallikrein).
(by kallikrein).
(by kallikrein).
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6 outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            outstation
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Length 621;

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Best Local (
PROTEIN SEQUENCE OF 387-455.

MEDLINE=76260155; PubMed=956151;
Han Y.N., Kato H., Iwanaga S., Suzuki T.;
"Primary structure of bovine plasma high-molecular-weight kininogen. The amino acid sequence of a glycopeptide portion (fragment 1) following the C-terminus ot the bradykinin moiety.";
J. Biochem. 79:1201-1222(1976).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KNG2 BOVIN STANDARD; PRT; 619 AA.
P0.1045; P0.1047;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
Kininogen-2 precursor (Kininogen II) (Thiol proteinase inhibitor)
[Contains: Kininogen-2 heavy chain; Bradykinin (Kallidin I); Lysy
bradykinin (Kallidin II); Kininogen-2 light chain].
                                                                                                                                                                                                                                    MEDLINE=70180420; PubMed=4986212;
Kato H., Nagasawa S., Suzuki T.;
"Studies on the structure of bovine kininogen: obonds and of methionyl bonds in kininogen-II.";
J. Biochem. 67:313-323(1970).
                                                                                                                                                                                                                                                                                                                                                                                                                                                Miyata T., Iwanaga S.;
"Bovine high molecular weight kiningen.
positions of carbohydrate chains and dischain portion.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    weight
Nature
[2]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nawa H., Kitamura N., Hirose T., Asai M., Inayama S., Nakanishi S.; "Primary structures of bovine liver low molecular weight kininogen precursors and their two mRNAB."; Proc. Natl. Acad. Sci. U.S.A. 80:90-94(1983).
                                                                                                                                                                                                                                                                                                                                           PROTEIN SEQUENCE (
MEDLINE=70180420;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUCLEOTIDE SEQUENCE [MRNA] (ISOFO MEDLINE=83117859; PubMed=6572010; Nawa H., Kitamura N., Hirose T.,
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MEDLINE=84014106; PubMed=6571699;
Kitamura N. Takagaki Y., Furuto S., Tanaka T., Nawa H., Nakanishi S.;
"A single gene for bovine high molecular weight and low molecular
"A single gene for bovine high molecular weight and low molecular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota, Metazoa,
Mammalia, Eutheria,
Pecora, Bovidae, Bov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BOVIN
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305:545-549(1983).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               124
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                                                                                                                                                                                                                                                                                                                                                                      OF 376-391
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Pred. No. 4.6e-36;
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PIR; A01282; KGBOH2.
PIR; A01284; KGBOL2.
HSSP; P01038; L390.
InterPro; IPR002395; Kininoge interPro; IPR00010; Prot interPro; PR00031; Cystatin; 3.
PRINTS; PR00331; Cystatin; 3.
PRINTS; SM00043; CY; 3.
PEPTIDE
PEPTIDE
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                                                                                                                           Vasoactive;
SIGNAL
                                                                                                                                         PROSITE; PS00287; CYSTATIN; 2.
Alternative splicing; Blood coagulation; Bradykinin; Direct protein sequencing; Glycoprotein; Hydroxylation; Inflammatory response; Plasma; Protease inhibitor; Pyrrolidone carboxylic acid; Repeat; Signal; Thiol protysoactive; Vasoactive;
                                                                                                                                                                                                                                                                                                                                EMBL;
EMBL;
PIR; A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HWW-kininogen plays an important role in blood coagulation by helping to position optimally prekallikrein and factor XI next to factor XII; (3) HWW-kininogen inhibits the thrombin-and plasmin-induced aggregation of thrombocytes; (4) the active peptide bradykinin that is released from HWW-kininogen shows a variety of physiological effects: (4A) influence in smooth muscle contraction, (4B) induction of hypotension, (4C) natriuresis and diuresis, (4D) decrease in blood glucose level, (4E) it is a mediator of inflammation and causes (4E1) increase in vascular permeability, (4E2) stimulation of nociceptors (4E3) release of other mediators of inflammation (e.g. prostaglandins), (4F) it has a cardioprotective effect (directly via bradykinin action, indirectly via endothellum-derived relaxing factor action); (5) LMW-kininogen inhibits the aggregation of thrombocytes; (6) LMW-kininogen is in contrast to HMW-kininogen not involved in blood
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROTEIN SEQUENCE OF 456-496.
MEDLINE-75170265; PubMed-1169237;
Han Y.N., Komiya M., Iwanaga S., Suzuki T.;
"Studies on the primary structure of bovine h
kininogen. Amino acid sequence of a fragment
peptide') released by plasma kallikrein.";
J. Biochem. 77:55-68(1975).
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    SUBCELLÜLAR LOCATIC
    ALTERNATIVE PRODUCT
    Event=Alternative &
    Name=HMW;

                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IsoId=P01045-2; Sequence=VSP_013564, VSP_013565; TISSUE SPECIFICITY: Plasma.
PTM: Bradykinin is released from kininogen by plasma kallikrein.
SIMILARITY: Contains 3 cystatin-like domains.
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V00427; CAA23710.1;
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H., Sugo T., Ikari
, Hotzer H. (eds.);
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Lysyl-bradykinin.
Bradykinin.
Kininogen-2 light
Cystatin-like 1.
Cystatin-like 2.
Cystatin-like 2.
Cystatin-like 3.
Not glycosylated.
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Kininogen-2 heavy
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Q6S911;
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05-JUL-2004 (TrEMBLrel. 2
01-FEB-2005 (TrEMBLrel. 2
01-FEB-2005 (TrEMBLrel. 2
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27, Last sequence update)
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29, Last annotation update)
(High molecular weight kininogen
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Cleavage (by kallikrein).
Cleavage (by kallikrein).
Pyrrolidone carboxylic acid.
4-hydroxyproline (By similarity).
N-linked (GlNAc. ); partial.
N-linked (GlNAc. ); partial.
N-linked (GlCNAc. );
N-linked (GlCNAc. );
N-linked (GlCNAc. );
N-linked (GlCNAc. );
O-linked (GlCNAc. )
O-linked (GlNAc. )
O-linked (GalNAc. )
                                                                                                                                                                                                                                                                                                                                                                                                                 isoform LMW).
/FTId=VSP_013564.
Missing (in isoform )
/FTId=VSP_013565.
T -> P.
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Pred. No. 1.6e-33;
4; Mismatches 22
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HVKSCEYKGRPQEAGAEPAPQGEVSLPAESPQLAR
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SQUERRE REPRESENTATION OF THE PROPERTY OF THE 
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RO 08677; 008676; Q91XK5;
DT 16-CCT-2001 (Rel. 40, Created)
DT 16-CCT-2001 (Rel. 40, Last sequence update)
DT 13-SEP-2005 (Rel. 48, Last annotation update)
DT 13-SEP-2005 (Rel. 48, Last annotation update)
DE Kininogen-1 precursor [Contains: Kininogen-1 heavy chain; Bradykinin;
DE Kininogen-1 light chain].
DE Kininogen-1 light chain]
DE Kininogen-1 Synonyms-Kng;
DE Kininogen-1 Nouse).
DE Kininogen-1 Nouse).
DE Kininogen-1 Light chain].
DE Light chain].
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GO; GO:0005515; C:extracellular space; GO; GO:0005515; C:extracellular space;
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Mammalia; Eutheria; Eu
Muridae; Murinae; Mus.
NCBI_TaxID=10090;
NUCLEOTIDE SEQUENCE [LARGE SCALE MRNA] (ISOFORM LMW).
STRAIN=CS7BL/60; TISSUE=Placenta;
MEDLINE=22354683; PubMed=12466851; DOI=10.1038/nature01266;
Okazaki Y., Furuno M., Kasukawa T., Adachi J., Bono H., Kon
Nikaido I., Osato N., Saito R., Suzuki H., Yamanaka I., Kiy
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MEDLINE=97342556; PubMed=9199253; DOI=10.1016/S0167-4781(97)00018-3;
Takano M., Kondo J., Yayama K., Otani M., Sano K., Okamoto H.;
Takano M., Kondo J., Yayama K., Otani M., Sano K., Okamoto H.;
"Molecular cloning of cDNAma for mouse low-molecular-weight and high-
molecular weight prekininogens.";
Biochim. Biophys. Acta 1352:222-230(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                               NUCLEOTIDE SEQUENCE [MRNA] (ISOFORMS STRAIN=C57BL/6 X CBA; TISSUE=Liver;
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to the EMBL/GenBank/DDBJ
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, Meneton P., Pesquero J.B.,
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RA Yagi K., Tomaru Y., Hasegawa Y., Nogami A., Schonbach C., Gojobori T.,
RA Baldarelli R., Hill D.P., Bult C., Hume D.A., Quackenbush J.,
RA Bchriml L.M.; Kanapin A., Matsuda H., Batalov S., Beisel K.W.,
RA Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA Blake J.A., Bradt D., Brusic V., Chothia C., Corbani L.E., Cousins S.,
RA Blake J.A., Bradt D., Brusic V., Godzik A., Grazer K.S.,
RA Blake J.A., Gariboldi M., Gissi C., Godzik A., Grazer K.S.,
RA Grimmond S., Gustincich S., Hirokawa N., Jackson I.J., Jarvis E.D.,
RA Kanai A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
RA Konagaya A., Kurochkin I.V., Lee Y., Lenhard B., Lyons P.A.,
RA Maglott D.R., Maltais L., Marchionni L., McKenzie L., Miki H.,
RA Nagashima T., Numata K., Okido T., Pavan W.J., Pertea G., Pesole G.,
RA Petrovsky N., Pillai R., Pontius J.U., Qi D., Ramachandran S.,
RA Pavasi T., Reed J.C., Reed D.J., Reid J., Ringwald M.,
RA Sultana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M.,
RA Sultana R., Takenaka Y., Taylor M.S., Teasdale R.D., Tomita M.,
RA Verardo R., Wagner L., Wahlestedt C., Wang Y., Watanabe Y., Wells C.,
RA Wilming L.G., Wynshaw-Boris A., Yanagisawa M., Yang I., Yang L.,
RA Himing L.G., Wynshaw-Boris A., Yanagisawa M., Yang I., Yang L.,
RA Himing L.G., Wynshaw-Boris A., Yanagisawa M., Sakazume N., Sato K.,
RA Himing L.G., Washizawa Y., Sakai K., Sasai K., Shibata K., Arakawa T., Pukuda S.,
RA Hasa A., Hashizume W., Imotani K., Jahit Y., Itoh M., Kagawa I.,
RA Yasunishi A., Yoshino M., Waterston R., Lander E.S., Rogers J.,
RA Yasunishi A., Hashizume W., Imotani K., Jahit Y., Itoh M., Kagawa I.,
RA Hasa A., Hashizume W., Imotani K., Jahit Y., Itoh M., Kagawa I.,
RA Hasanishi A., Koshino M., Waterston R., Lander E.S., Rogers J.,
RA Yasunishi A., Sakai K., Kawai J., Arakawa T., Shinasa Y.,
RA Yasunishi A., Sakai K., Kanasi J., Ra Waterston R., Lander E.S., Rogers J.,
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"Analysis of the mouse transcriptome
60,770 full-length cDNAs.";
Nature 420:563-573(2002).
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                               Event-Alternative Name-HMW;
IsoId=008677-1; Sequence=Displayed;
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                                                        splicing; Named isoforms=2
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InterPro; IPR000010; Prot inh_cystat
of=m. pr00031; Cystatin; 3.
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EMBL;
EMBL;
                                                                                                                                                                             SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This Swiss-Prot entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Ew the European Bioinformatics Institute. There are no restr
                                                                                                                                                                                                    VARSPLIC
                                                                                                                                                                                                                                                                                                                                       DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRINTS; PR00334; KININOGEN PROSITE; PS00287; CYSTATIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GO; GO:0005615; C:extracellular space;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IsoId=008677-2; Sequence=VSP_001263, VSP_001264; TISSUE SPECIFICITY: Plasma.
FTM: Bradykinin is released from kininogen by plasma kallikrein.
SIMILARITY: Contains 3 cystatin-like domains.
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                                          311
                                                                                      251
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D84415; BAA19742.1; -; mRNA.
AKO05547; -; NOT ANNOTATED CDS; mRNA.
BC018158; AAH18158.1; -; mRNA.
                                                               62
                                                                                                            N
                                                                                                                                  82;
                                                                                                                                             Similarity
ALDM 374
                   PLGM 125
                                          VAGTKYVIEFIARETKCSKESNTELAEDCEIKHLGQSLDCNANYMRPWENKVVPTVKCQ
                                                                VAGKKYFIDFVARETTCSKESNEELTESCETKKLGQSLDCNAEVYVVPWEKKIYPTVNCQ
                                                                                      SGDDLVEALPKPCPGCPRDIPVDSPELKEVLGHSIAQLNAENDHPFYYKIDTVKKATSQV
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By similarity.
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N-linked (GlcNAc. . .) (Pot
N-linked (GlcNAc. . .) (Pot
N-linked (GlcNAc. . .) (Pot
Interchain (between heavy a
chains) (By similarity).
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Cystatin-like 1.
Cystatin-like 2.
Cystatin-like 3.
His-rich.
                                                                                                                                             Pred.
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/FTId=VSP
                                                                                                                                                       Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bradykinin.
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RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausmer R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
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RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Fahey J., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
RA Butterfield Y.S.N., Krzywinski M.I., Skalaka U., Smallus D.E.,
"Generation and initial analysis of more than 15,000 full-length human
"Generation and initial analysis of more than 15,000 full-length human
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KNG1 RAT STANDARI
P08934; P08933;
01-NOV-1988 (Rel. 09, 0
01-NOV-1988 (Rel. 09, 1
13-SEP-2005 (Rel. 48, 1
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01-FEB-2005
01-FEB-2005
01-FEB-2005
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Submitted (DEC-2004) to the EMBL/GenBank/DDBJ databases.
Submit ed (DEC-2004) to the EMBL/GenBank/DDBJ databases.
SMBL; BC088155; AAH88155.1; -; mRNA.
GO; GO:0004869; F:cysteine protease inhibitor activity; IEA.
GO; GO:0004869; F:cysteine protease inhibitor activity; IEA.
SEQUENCE 433 AA; 47891 MW; 6D584348595166ED CRC64;
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
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llarity 65.0%;
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Pred. No. 1.9e
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J. Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                       Furuto-Kato S., Matsumoto A., Kitamura N., Nakanishi S.; Primary structures of the mRNAs encoding the rat precursors for bradykinin and T-kinin. Structural relationship of kininogens with major acute phase protein and alpha 1-cysteine proteinase inhibitor."; J. Biol. Chem. 260:12054-12059(1985).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  family."
J. Biol.
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NUCLEOTIDE SEQUENCE [MRNA] (ISOFORMS HMW AND LMW).

MEDLINE=87137443; PubMed=3029068;

Kitagawa H., Kitamura N., Hayashida H., Miyata T., Nakanishi S.;

Kitagawa H., Kitamura N., Hayashida H., Miyata T., Nakanishi S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognathi;
Muroidea; Muridae; Murinae; Rattus.
NCBI_TaxID=10116;
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Kininogen-1 light chain].
Name=Kng1; Synonyms=Kng;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kageyama R., Kitamura N., Ohkubo H., Nakanishi S.;
"Differing utilization of homologous transcription initiatirat K and T kininogen genes under inflammation condition.",
J. Biol. Chem. 262:3345-2351(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Fung W.-P., Schreiber G.; "Structure and expression of the genes for major acute phase alpha protein (thiostatin) and kininogen in the rat."; J. Biol. Chem. 262:9298-9308(1987).
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between
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=86008264; PubMed=2413018;
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                                                                                                                                                                                                                                                                                                                          SUBCELLULAR LOCATION:
ALTERNATIVE PRODUCTS:
Event=Alternative spl
                                                                                                                           PTM: Bradykinin is released from kininogen by plasma kallikrein.
MISCELLANEOUS: Rats express four types of kininogens: the
classical HMW/LMW kininogens and two additional LMW-like
                                                                        SIMILARITY: Contains 3 cystatin-like domains.
                                                                                                    classical HMW/LMW kininogens and kininogens: T-I and T-II.
                                                                                                                                                                                                      TISSUE SPECIFICITY:
                                                                                                                                                                                                                                                                                                                                                                                                     clotting.
Swiss-Prot entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                IsoId=P08934-2;
                                                                                                                                                                                                                                                                               IsoId=P08934-1;
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                                                                                                                                                                                                                           Sequence=VSP_001265,
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                                                                                                                                                                                                                                                                                                                                                                                 Secreted;
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gous transcription initiation sites
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Query Match
Best Local S
Matches 80
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EMBL; M11884; AAA41887.1;
EMBL; M14369; AAA41885.1; A
EMBL; M14369; AAA41485.1; A
EMBL; M14369; AAA41482.1; A
EMBL; M16455; AAA41482.1; APER; A25486; A25486.
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CARBOHYD
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ProDom; PD001231; Prot inh I25A_B; 1.

PROSITE; PS00287; CYSTĀTIN; 2.

Alternative splicing; Blood coagulation;

Inflammatory response; Multigene family;

Repeat; Signal; Thiol protease inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HSSP; P01040; IDVD.

RGD; 2980; kngl.
InterPro; IPR0002395; Kininogen.
InterPro; IPR000010; Prot_inh_cystat.
InterPro; IPR003243; Prot_inh_I25A_B.
Pfam; PP00031; Cystatin; 3.
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By similarity.

CEYKGRLIKAGAGPAPERQAEASTVTP (in isoform
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/FTIG=VSP 001265.
Missing (In isoform LMW).
/FTId=VSP 001266.
E -> K (in Ref. 2).
B -> K (in Ref. 2).
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Pred. No. 2.9e
14; Mismatches
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Kininogen-1 light of Cystatin-like 1.
Cystatin-like 2.
Cystatin-like 3.
His-rich.
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MGI; MGI:3027157; Kng2.
GO; GO:0004869; F:Cysteine protease inh
InterPro; IPR000010; Prot_inh_Cystat.
Pfam; PF00031; Cystatin; 3.
SMART; SM00043; CY; 3.
PROSITE; PS00289; CYSTATIN; 1.
SEQUENCE 433 AA; 47887 MW; C86F7BE9
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Q6S912;
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05-JUL-2004
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"Structure and expression of two kir Biol. Chem. 385:295-301(2004).
EMBL; AY462057; AAR88631.1; -; mRNA. MGI; MGI:3027157; Kng2.
GO; GO:0004869; F:cysteine protease
                                                                                                                                                                                     STRAIN=FVB/N;
MEDLINE=23306569; PubMed=15134344; DOI=10.1515/BC.2004.025;
Cardoso C.C., Garrett T., Cayla C., Meneton P., Pesquero J.
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Mammalia; Eutheria; Eu
Muridae; Murinae; Mus.
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MEDLINE=23306569; PubMed=15134344; DOI=10.1515/BC.2004.025;
Cardoso C.C., Garrett T., Cayla C., Meneton P., Pesquero J.B.,
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Q6S910 MOUSE PRELIMINARY;
Q6S910;
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Muridae; Murinae; Mus.
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Pred. No. 2.4e-31;
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pfam; PP00031; Cystatin; 3.
SMART; SM00043; CY; 3.
PROSITE; PS00287; CYSTATIN; 1.
SEQUENCE 462 AA; 51052 MW;
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Mammalia; Eutheria; Euarchontoglires; Glires; Rodent
Muridae; Murinae; Mus.
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05-JUL-2004 (TREMBLrel.
05-JUL-2004 (TREMBLrel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Structure and expression of two kininogen Biol. Chem. 385:295-301(2004).
EMBL, AY462056; AAR88630.1; -; mRNA.
MGI; MGI:3027157; Kng2.
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SMART; SM00043; CY; 3.
PROSITE; PS00287; CYSTATIN;
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Rodentia; Sciurogna
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13-SEP-2005 (Rel. 48, Last sequence update)
13-SEP-2005 (Rel. 48, Last annotation update)
T-kininogen II precursor (Majora caute phase protein) (Alpha-1-MAP)
(Thiostatin) [Contains: T-kininogen II heavy chain; T-kininogen II light chain].
kininogen II light chain].
Rattus norvegicus (Ret).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Sciurognat
Maroidea; Muridae; Murinae; Rattus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          kininogens.";
J. Biol. Chem.
                                                                                                                                                                                                               EMBL; M11885; AAA41491.1; -; mRNA.
EMBL; M11661; AAA41570.1; ALT_FRAME; mRNA.
PIR; B28055; B28055; B28055; B28055; B28055; -..
GlycoSulteDB; F08932; -..
Ensembl; ENSRNOG00000030387; Rattus norvegicus.
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InterPro; IPR000010; Prot_inh_cystat.
InterPro; IPR003243; Prot_inh_I25A_B:
Pfam; PF00031; Cystatin; 3
Probom; PD001231; Prot_inh_I25A_B; 1.
SMART; SM00043; CY; 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This Swiss-Prot entry is copyright. It is produced through a detween the Swiss Institute of Bioinformatics and the EMBL the European Bioinformatics Institute. There are no restrict use as long as its content is in no way modified and this statuse as long as its content is in no way modified and this statuse as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as its content is in no way modified and this status as long as long as its content is in no way modified and this status as long as long as its content is in no way modified and this status as long as long as its content is in no way modified and this status as long as long as its content is in no way modified and this status as long as 
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SUBCELLULAR LOCATION: Extra
TISSUE SPECIFICITY: Plasma.
INDUCTION: In response to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: Kininogens are plasma glycoproteins with a number of functions: (1) as precursor of the active peptide bradykinin they effect smooth muscle contraction, induction of hypotension and increase of vascular permeability. (2) They play a role in blood coagulation by helping to position optimally prekallikrein and factor XI next to factor XII. (3) They are inhibitor of thiol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the T-II.
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PTM: As T kinin is preceded by a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MISCELLANEOUS: Rats classical HMW and LA
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; ALT_FRAME;
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PROSITE; PS00287; CYSTATIN; 2.
Acute phase; Bradykinin; Direct protein sequencing; Glycoprotein;
Multigene family; Plasma; Protease inhibitor; Repeat; Signal;
Thiol protease inhibitor; Vasoactive; Vasodilator.

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MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins P.S., Wagner L., Shenmen C.M., Schuler
                                                                                                                                                                                                                                                                   Similar to alpha-1 major
Name=MGC108747;
Rattus norvegicus (Rat).
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                                                                                                             NUCLEOTIDE SEQUENCE.
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ALISCHUI S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
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Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
"Generation and initial analysis of more than 15,000 full-length human transmission of the start 
      RESULT 15
Q7YRP6_PIG
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Director MGC Project;
Submitted (DEC-2004) to the EMBL/GenBank/DDBJ databases.
SMBL; BC088161; AAH88161.1; -; mRNA.
GO; GO:0004869; F:cysteine protease inhibitor activity; IEA.
GO; GO:0004869; F:cysteine protease.
InterPro; IPR000010; Prot_inh_Cystat.
InterPro; IPR003243; Prot_inh_I25A_B.
InterPro; IPR003243; Prot_inh_I25A_B.
InterPro; IPR003243; Prot_inh_I25A_B.
InterPro; IPR003243; Prot_inh_I25A_B.
NUCLEOTIDE SEQUENCE.

PubMed=13679312; DOI=10.1095/biolreprod.103.020412;

Vonnahme K.A., Fernando S.C., Ross J.W., Ashworth M.D.,

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Malayer J.R., Geisert R.D.;

Porcine endometrial expression of kininogen, factor XII

"Porcine endometrial expression of gilts.";

Biol. Reprod. 70:132-138(2004).

EMBL; AY321363; AAP85260.1; -; mRNA.
                                                                                                                                                                                                                                                                   Name=kng;
Sus scrofa (Pig)
Eukaryota; Metazo
                                                                                                                                                                                                                                                                                                                               01-OCT 2003 (TrEMBLrel. 25, Created)
01-OCT-2003 (TrEMBLrel. 25, Last sequence update)
01-WAR-2004 (TrEMBLrel. 26, Last annotation update)
Low molecular weight kininogen (Fragment).
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ProDom; PD001231; Prot_inh I25A B; 1.
SMART; SM00043; CY; 3.
PROSITE; PS00287; CYSTATIN; 2.
                                                                                                                                                                                                  NCBI_TaxID=9823;
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Director MGC
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nilarity 61.8%;
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